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(54) Title: METHOD OF INHIBITING CATHEPSIN K



Human Cathepsin K

#### (57) Abstract

A nevel cathepsin K crystalline structure is identified. Also disclosed are methods of identifying inhibitors of this protease and methods of inhibiting cathepsin K using inhibitors with certain structural, physical and spatial characteristics.

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#### METHOD OF INHIBITING CATHEPSIN K

#### Field of the Invention

This invention relates to a method of inhibiting cathepsin K by administering compounds with certain structural, physical and spatial characteristics that allow for the interaction of said compounds with specific residues of the active site of the enzyme. This interaction between the compounds of this invention and the active site inhibits the activity of cathepsin K and these compounds are useful for treating diseases in which said inhibition is indicated, such as osteoporosis and periodontal disease. This invention also relates to a novel crystalline structure of cathepsin K, the identification of a novel protease catalytic active site for this enzyme and methods enabling the design and selection of inhibitors of said active site.

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#### Background of the Invention

Cathepsin K is a member of the family of enzymes which are part of the papain superfamily of cysteine proteases. Cathepsins B, H, L, N and S have been described in the literature. Recently, cathepsin K polypeptide and the cDNA encoding such polypeptide were disclosed in U.S. Patent No. 5,501,969 (called cathepsin O therein). Cathepsin K has been recently expressed, purified, and characterized. Bossard, M. J., et al., (1996) J. Biol. Chem. 271, 12517-12524; Drake, F.H., et al., (1996) J. Biol. Chem. 271, 12511-12516; Bromme, D., et al., (1996) J. Biol. Chem. 271, 2126-2132.

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Cathepsin K has been variously denoted as cathepsin O, cathepsin X or cathepsin O2 in the literature. The designation cathepsin K is considered to be the more appropriate one (name assigned by Nomenclature Committee of the International Union of Biochemistry and Molecular Biology).

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Cathepsins of the papain superfamily of cysteine proteases function in the normal physiological process of protein degradation in animals, including humans, e.g., in the degradation of connective tissue. However, elevated levels of these enzymes in the body can result in pathological conditions leading to disease. Thus, cathepsins have been implicated in various disease states, including but not limited to, infections by pneumocystis carinii, trypsanoma cruzi, trypsanoma brucei brucei, and Crithidia fusiculata; as well as in schistosomiasis malaria, tumor metastasis, metachromatic leukodystrophy, muscular dystrophy, amytrophy, and the like. See International Publication Number WO 94/04172, published on March 3, 1994, and

references cited therein. See also European Patent Application EP 0 603 873 A1, and references cited therein. Two bacterial cysteine proteases from P. gingivallis, called gingipains, have been implicated in the pathogenesis of gingivitis. Potempa, J., et al. (1994) Perspectives in Drug Discovery and Design, 2, 445-458.

Cathepsin K is believed to play a causative role in diseases of excessive bone or cartilage loss. Bone is composed of a protein matrix in which spindle- or plate-shaped crystals of hydroxyapatite are incorporated. Type I Collagen represents the major structural protein of bone comprising approximately 90% of the structural protein. The remaining 10% of matrix is composed of a number of non-collagenous proteins, including osteocalcin, proteoglycans, osteopontin, osteonectin, thrombospondin, fibronectin, and bone sialoprotein. Skeletal bone undergoes remodeling at discrete foci throughout life. These foci, or remodeling units, undergo a cycle consisting of a bone resorption phase followed by a phase of bone replacement.

Bone resorption is carried out by osteoclasts, which are multinuclear cells of hematopoietic lineage. The osteoclasts adhere to the bone surface and form a tight sealing zone, followed by extensive membrane ruffling on their apical (i.e., resorbing) surface. This creates an enclosed extracellular compartment on the bone surface that is acidified by proton pumps in the ruffled membrane, and into which the osteoclast secretes proteolytic enzymes. The low pH of the compartment dissolves hydroxyapatite crystals at the bone surface, while the proteolytic enzymes digest the protein matrix. In this way, a resorption lacuna, or pit, is formed. At the end of this phase of the cycle, osteoblasts lay down a new protein matrix that is subsequently mineralized. In several disease states, such as osteoporosis and Paget's disease, the normal balance between bone resorption and formation is disrupted, and there is a net loss of bone at each cycle. Ultimately, this leads to weakening of the bone and may result in increased fracture risk with minimal trauma.

The abundant selective expression of cathepsin K in osteoclasts strongly suggests that this enzyme is essential for bone resorption. Thus, selective inhibition of cathepsin K may provide an effective treatment for diseases of excessive bone loss, including, but not limited to, esteoporosis, gingival diseases such as gingivitis and periodontitis. Paget's disease, hypercalcemia of malignancy, and metabolic bone disease. Cathepsin K levels have also been demonstrated to be elevated in chondroclasts of osteoarthritic synovium. Thus, selective inhibition of cathepsin K may also be useful for treating diseases of excessive cartilage or matrix degradation,

including, but not limited to, osteoarthritis and rheumatoid arthritis. Metastatic neoplastic cells also typically express high levels of proteolytic enzymes that degrade the surrounding matrix. Thus, selective inhibition of cathepsin K may also be useful for treating certain neoplastic diseases.

Surprisingly, it has been found that a broad, structurally diverse series of compounds have common structural, physical and spatial characteristics that allow for the interaction of said compounds with specific residues of the active site of cathepsin K and are useful for treating diseases in which inhibition of bone resorption is indicated, such as osteoporosis and periodontal disease. Thus, this invention relates to the method of inhibiting cathepsin K using compounds having the characteristics hereinbelow defined.

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#### Summary of the Invention

In one aspect, the present invention provides a method for inhibiting cathepsin K by administering compounds with certain structural, physical and spatial characteristics that allow for the interaction of said compounds with specific residues of the active site of the enzyme. This interaction inhibits the activity of cathepsin K and, thus, treats diseases in which bone resorption is a factor.

In another aspect, the present invention provides a novel cysteine protease in crystalline form.

In yet another aspect, the invention provides a novel protease composition characterized by a three dimensional catalytic site formed by the atoms of the amino acid residues listed in Table XXIX.

In still another aspect, the invention provides a method for identifying inhibitors of the compositions described above which methods involve the steps of: providing the coordinates of the protease structure of the invention to a computerized modeling system; identifying compounds which will bind to the structure; and screening the compounds or analogs derived therefrom identified for cathepsin K inhibitory bioactivity.

Other aspects and advantages of the present invention are described further in the following detailed description of the preferred embodiments thereof.

#### Brief Description of the Drawings

Figure 1 is the amino acid sequence of cathepsin K aligned with the amino acid sequences of other cysteine proteases.

Figure 2 is a ribbon diagram of cathepsin K. The amino and carboxyltermini are indicated by N and C. The drawing was produced using the program MOLSCRIPT [Kraulis, P., J. Appl. Crystallogr., 24, 946-950 (1991)].

Figure 3 is a ribbon diagram of cathepsin K in complex with E-64, a known inhibitor of cysteine proteases. The drawing was produced using the program MOLSCRIPT.

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Figure 4a is an illustration of the active site of cathepsin K. Figure 4b is a stereoview of the active site of cathepsin K. For clarity, no hydrogen atoms or water molecules are shown.

Figures 5a-13a are illustrations of the active site of cathepsin K in complex with novel inhibitors of cathepsin K. Figures 5b-13b are stereoviews of the active site of cathepsin K in complex with novel inhibitors of cathepsin K. These views depict the interaction of each inhibitor with all atoms of residues of the active site of cathepsin K within 5Å of the inhibitors. For clarity, no hydrogen atoms or water molecules are shown.

Table I provides the three dimensional protein coordinates of the cathepsin K crystalline structure of the invention.

Tables II-X provide the three dimensional coordinates for the cathepsin K complex with specific inhibitors of the present invention.

Tables XI-XIX provide listings of the three atom angles between atoms of the inhibitors and the protein for all inhibitor atoms within 5 Ångstroms of the protein.

Tables XX-XXVIII provide listings of the distances between atoms of the inhibitors and the protein for all inhibitor atoms within 5 Ångstroms of the protein.

Table XXIX provides the atoms of the amino acid residues of the catalytic site.

#### **Detailed Description of the Invention**

The present invention provides a novel cysteine protease crystalline structure, a novel cysteine protease active site, and methods of use of the crystalline form and active site to identify protease inhibitor compounds.

In particular, the present invention provides a method for inhibiting cathepsin K by administering compounds with certain structural, physical and spatial characteristics that allow for the interaction of said compounds with specific residues

Specifically, the inhibitors of cathepsin K used in the present invention interact with any two or more of the following:

- 1. Tyrosine 67 sidechain;
- 2. Hydrophobic pocket lined with atoms from methinoine 68,
- 5 leucine 209, alanine 163, alanine 134 and portions of tyrosine 67;
  - 3. Hydrogen bonds donated by glycine 66 amide nitrogen;
  - 4. Cysteine 25 the active site nucleophile;
  - 5. Mainchain interactions from residues glutamine 21, cysteine 22, and glycine 23;
- 10 6. Tryptophan 184 sidechain; and

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7. Hydrophobic contacts with the sidechain atoms of glutamine 143 and asparagine 161 and the mainchain of alanine 137 and serine 138.

Preferably, the inhibitors of cathepsin K used in the present invention interact with any three or more of the above-identified regions of the active site.

The compounds used in the methods of the present invention possess an electrophilic carbon and either a hydrophobic group whose centroid is 5.44-6.94Å from the carbon or an aromatic group whose centroid is 9.24-11.24Å from the carbon, or both the hydrophobic and the aromatic groups in which case the centroids of these two groups should be 15.67-16.67Å apart. These features must be able to make the appropriate interactions with the cathepsin K active site. The electrophilic carbon atom should be 1.7-4.0Å from the side chain sulfur atom (SG) on the amino acid cysteine 25. The hydrophobic group should be near the following amino acids with appropriate distance ranges between the centroid of the side chain atoms and the centroid of the hydrophobic group given in parentheses: tyrosine 67 (4.91-5.91Å), methionine 68 (5.74-6.74Å), alanine 134 (4.15-5.15Å), leucine 160 (6.18-7.18Å), and leucine 209 (5.71-6.71Å). The aromatic group should be near the either

The key structural features of the inhibitors of the present invention include an electrophilic carbon, preferably the carbon of a carbonyl group, a hydrophobic group, preferably an isobutyl group, and an aromatic group, preferably a phenyl group. The electrophilic carbon of the inhibitor may be in the same compound with two hydrophobic groups, such as two isobutyl groups, or two aromatic groups, such as two phenyl groups, or one hydrophobic group and one aromatic group.

Suitably, the method of inhibiting cathepsin K of the present invention comprises administering to a mammal, preferably a human, in need thereof a

tryptophan 184 (4.10-7.10Å) or tryptophan 188 (4.10-7.10Å) or both.

compound that fits spatially into the active site of cathepsin K, said compound comprising any two or more of the following:

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(i) an electrophilic carbon atom that binds to the side chain sulfur atom of cysteine 25 wherein said electrophilic carbon atom is 1.7-4.0Å from said sulfur atom;

- (ii) a hydrophobic group that interacts with tryptophan 184 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tryptophan 184 is 4.10-7.10Å;
- (iii) a hydrophobic group that interacts with tyrosine 67, methionine 68, alanine 134, leucine 160, and leucine 209, creating a hydrophobic pocket, and has distance ranges between the centroid of said hydrophobic group and the centroids of the side chain atoms of the amino acid residues of said hydrophobic pocket which are tyrosine 67: 4.91-5.91Å, methionine 68: 5.74-6.74Å, alanine 134: 4.15-5.15Å, leucine 160: 6.18-7.18Å, and leucine 209: 5.71-6.71Å;
- (iv) a hydrophobic group that interacts with tyrosine 67 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tyrosine 67 is 4.10-7.10Å;
- (v) an amino group with a pKa of less than 7 or an oxygen atom, each of which interacts with a hydrogen atom donated by the amide nitrogen of glycine 66 wherein the distance between these two atoms is 2.7-3.5Å;
- (vi) a hydrophobic group that interacts with the main chain atoms of glutamine 21, cysteine 22 and glycine 23 wherein the distance between the centroid of said hydrophobic group and the centroids of glutamine 21, cysteine 22 and glycine 23 are 3.7-5.4, 4.9-5.7 and 5.4-6.7Å, respectively; or
- (vii) a hydrophobic group that interacts with the side chain atoms of glutamine 143 and asparagine 161 and the main chain of alanine 137 and serine 138 wherein the distance between the centroid of the hydrophobic group and the centroids of glutamine 143, asparagine 161, alanine 137, and serine 138 are 7.9-9.6Å, 4.7-5.4Å, 4.2-5.5Å, and 4.6-6.4Å, respectively. Preferably, the inhibitors of cathepsin K used in the present invention comprise three or more of the above.

Suitably, the method of inhibiting cathepsin K of the present invention comprises administering to a mammal, preferably a human, in need thereof, a compound that fits spatially into the active site of cathepsin K, said compound comprising:

(i) an electrophilic carbon atom that binds to the side chain sulfur atom of cysteine 25 wherein said electrophilic carbon atom is 1.7-4.0Å from said sulfur atom; and

(ii) a hydrophobic group that interacts with tryptophan 184 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tryptophan 184 is 4.10-7.10Å. Preferably, the hydrophobic group that interacts with tryptophan 184 is an aromatic group and the centroid of this aromatic group is 9.24-11.24Å from the centroid of the electrophilic carbon that binds to the side chain sulfur atom of cysteine 25.

Preferably, the electrophilic carbon that binds to the side chain sulfur atom of cysteine 25 is a carbonyl carbon.

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Suitably, the method of the present invention further comprises a compound with a hydrophobic group that:

has a centroid which is 5.44-6.94Å from said electrophilic carbon; interacts with tyrosine 67, methionine 68, alanine 134, leucine 160, and leucine 209, creating a hydrophobic pocket; and

has distance ranges between the centroid of said hydrophobic group and the centroids of the side chain atoms of the amino acid residues of said hydrophobic pocket which are tyrosine 67: 4.91-5.91Å, methionine 68: 5.74-6.74Å, alanine 134: 4.15-5.15Å, leucine 160: 6.18-7.18Å, and leucine 209: 5.71-6.71Å. Preferably, this hydrophobic group is an isobutyl group.

Alternately, the method of the present invention further comprises a compound with a hydrophobic group that interacts with tyrosine 67 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tyrosine 67 is 4.10-7.10Å. Preferably, this hydrophobic group is an aromatic group.

Alternately, the method of the present invention further comprises a compound with an amino group with a pKa of less than 7 or an oxygen atom, each of which interacts with a hydrogen atom donated by the amide nitrogen of glycine 66 wherein the distance between these two atoms is 2.7-3.5Å. Preferably, the compound comprises an oxygen atom, such as an oxygen atom of a carbonyl group or an oxygen atom of a hydroxyl group.

Alternately, the method of the present invention further comprises a compound with a hydrophobic group that interacts with the main chain atoms of glutamine 21, cysteine 22 and glycine 23 wherein the distance between the centroid

of the hydrophobic group and the centroids of glutamine 21, cysteine 22 and glycine 23 are 3.7-5.4, 4.9-5.7 and 5.4-6.7Å, respectively. Preferably, this hydrophobic group is an isobutyl group.

Alternately, the method of the present invention further comprises a compound with a hydrophobic group that interacts with the side chain atoms of glutamine 143 and asparagine 161 and the mainchain of alanine 137 and serine 138 wherein the distance between the centroid of the hydrophobic group and the centroids of glutamine 143, asparagine 161, alanine 137, and serine 138 are 7.9-9.6Å, 4.7-5.4Å, 4.2-5.5Å, and 4.6-6.4Å, respectively.

Compounds used in the method of the present invention include, but are not limited to, the following:

3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2hexanone:

4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-

[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone; 15

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4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N-[N-(methyl)-L-leucyl)]-3pyrrolidinone;

4-[N-[(phenyimethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone;

bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one;

2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-Lleucinyl)]carbohydrazide:

(1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide;

1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)amino-propan-2-one; and

2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide; or a pharmaceutically acceptable salt thereof.

As stated herein, the interaction of the inhibitor at the side chain sulfur atom of cysteine 25 has as one of its requirements that the inhibitor contain an 30 "electrophilic carbon" atom. By this term is meant an electron deficient carbon. This term includes, but is not limited to, a carbonyl carbon atom. This term also includes an epoxide, a thiocarbonyl, an imine, and a nitrile. Suitably, this term may also be represented by the formula -C=N-X, wherein X may be optionally tied back to C in a ring or wherein X is CH2, H, O, S or NRa in which Ra is H of C1-4alkyl.

includes an epoxide, a thiocarbonyl, an imine, and a nitrile. Suitably, this term may also be represented by the formula -C=N-X, wherein X may be optionally tied back to C in a ring or wherein X is  $CH_2$ , H, O, S or  $NR^a$  in which  $R^a$  is H of  $C_{1-4}$  alkyl.

The hydrophobic groups that interact with tryptophan 184 or tyrosine 67 include, but are not limited to, aromatic groups. These hydrophobic groups include phenyl,  $C_{1-6}$ alkyl and heteroaryl, which is defined hereinbelow. The hydrophobic groups that interact with the hydrophobic pocket lined with atoms from tyrosine 67, methionine 68, alanine 134, leucine 160, and leucine 209 not only includes isobutyl, but also includes  $C_{1-6}$ alkyl,  $C_{3-6}$ cycloalkyl and adamantyl. The hydrophobic groups that interact with the main chain atoms of glutamine 21, cysteine 22 and glycine 23 or the side chain atoms of glutamine 143 and asparagine 161 and the mainchain of alanine 137 and serine 138 include  $C_{1-10}$ alkyl,  $C_{b}F_{2b+1}$ , in which b is 1-3, and aryl and heteroaryl, each of which are defined hereinbelow.

As used herein, the term "centroid" means the position for the stated atoms calculated by averaging the x coordinates of the atoms to obtain the x coordinate of the centroid, averaging the y coordinates of the atoms to obtain the y coordinate of the centroid, and averaging the z coordinates of the atoms to obtain the z coordinate of the centroid.

The compounds used in the method of the present invention include, but are not limited to, the compounds of formula (I):

wherein:

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Q=

where:

$$A = absent,$$
 $R^{B}$ 

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$$B = X = Y , O R^{15}$$

 $L = C_{2-6} \text{alkyl}, \text{Ar-}C_{0-6} \text{alkyl}, \text{Het-}C_{0-6} \text{alkyl}, \text{CH}(R^{66}) \text{NR}^{60} \text{R}^{68},$   $\text{CH}(R^{66}) \text{Ar}, \text{CH}(R^{66}) \text{OAr'}, \text{NR}^{66} \text{R}^{67};$ 

10  $M = C(O), SO_2;$ 

G =

 $J = C(O), SO_2;$ 

T = Ar, Het;

 $V = C_{3-7}$ cycloalkyl;

 $W = H, -CN, -CF_3, -NO_2, -COR^7, -CO_2R^6, -CONHR^6,$   $-SO_2NHR^6, -NHSO_2R^6, -NHCOR^7, -O-COR^6, -SR^6,$   $NR'R^6, NR'(C=NH)NHR^5, Cl, Br, I, F;$ 

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X = Y = Z = N, O, S or CR^4,
                                                                                               provided that at least two of X, Y and Z are heteroatoms
                                                                                               and at least one of X, Y and Z is N, or one of X, Y and Z is
                                                                                               C=N, C=C or N=N and the other two are CR4 or N,
                                                                                               provided that X, Y and Z together comprise at least two N;
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                                                                      = indicates a single or double bond in the five-membered
                                                                       heterocycle;
                                                                      m = 0, 1, 2;
                                                                      n = 1 \text{ to } 6;
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                                                                      f = 0, 1, 2;
                                                                     Ar = phenyl, naphthyl, optionally substituted by one or more of
                                                                                             Ph-C_{0-6}alkyl, Het-C_{0-6}alkyl, C_{1-6}alkoxy, Ph-C_{0-6}alkoxy,
                                                                                             Het-C<sub>0-6</sub>alkoxy, OH, (CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>,
                                                                                             O(CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>:
                                                                     Ar' = phenyl or naphthyl, optionally substituted by one or more of
    15
                                                                                            \label{eq:condition} {\it Ph-C_{0-6}alkyl,\, Het-C_{0-6}alkyl,\, C_{1-6}alkoxy,\, Ph-C_{0-6}alkoxy,\, C_{1-6}alkoxy,\, C_{1-6}alkox
                                                                                           Het-C<sub>0-6</sub>alkoxy, OH, (CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>,
                                                                                           O(CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>, or halogen;
                                                                    R' = H, C_{1-6}alkyl, Ar-C_{0-6}alkyl, Het-C_{0-6}alkyl;
                                                                   R^! = H, C_{1-6alkyl};
  20
                                                                  R<sup>2</sup> = C<sub>4</sub>-6alkyl, C<sub>4</sub>-6alkenyl, benzyl;
                                                                  R^3 = C_{1-6}alkyl, Ar-C_{0-6}alkyl, Het-C_{0-6}alkyl, R<sup>5</sup>CO-, R<sup>5</sup>SO<sub>2</sub>-,
                                                                                          R5OC(O)-, R5NHCO-;
                                                                  R^4 = H, C_{1-6}aikyl, A_{1-C_{1-6}}alkyl, Het-C_{0-6}alkyl;
                                                                 R^5 = Ar-0-6alkyl, Het-C_{0-6}alkyl;
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                                                                 R^6 = H, C_{1-6aikyl}, CH_2CF_3, Ar-C_{0-6aikyl}, Het-C_{0-6aikyl};
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 $R^7 = C_{1-6}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

 $R^8$  = H; C<sub>2-6</sub> alkenyl; C<sub>2-6</sub>alkynyl; Het; Ar; C<sub>1-6</sub>alkyl, optionally substituted by OR', SR', NR'<sub>2</sub>, CO<sub>2</sub>R', CO<sub>2</sub>NR'<sub>2</sub>, N(C=NH)NH<sub>2</sub>, Het or Ar;

 $R^9 = H$ ,  $C_{1-6}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

 $R^{10} = C_{1-6}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

 $R^{11}$  = H,  $C_{1-6}$ alkyl, Ar- $C_{1-6}$ alkyl, Het- $C_{0-6}$ alkyl, or

$$R^{17}R^9-N$$

 $R^{12} = H, C_{1-6}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

 $R^{13} = H$ ,  $C_{1-6}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

$$R^{14} = \frac{R^{19}}{N - R^9 R^{72}, Ac}$$

 $R^{15} = H$ ,  $C_{1-6}$ alkyl,  $C_{2-6}$ alkenyl,  $C_{2-6}$ alkynyl, Ar, Het, or  $C_{1-6}$ alkyl optionally substituted by  $OR^9$ ,  $NR^9_2$ ,  $CONR^9_2$ , N(C=NH)NH-, Het or Ar;

 $R^{16} = C_{2-6}$ alkyl,  $C_{2-6}$ alkenyl,  $C_{2-6}$ alkynyl, Ar, Het, or  $C_{2-6}$ alkyl optionally substituted by  $OR^9$ ,  $SR^9$ ,  $NR^9_2$ ,  $CO_2R^9$ ,  $CONR^9_2$ , N(C=NH)NH-, Het or Ar;

 $R^{19} = H$ ,  $C_{1-6}$ alkyl,  $C_{2-6}$ alkenyl,  $C_{2-6}$ alkynyl, Ar, Het, or  $C_{1-6}$ alkyl optionally substituted by  $OR^9$ ,  $SR^9$ ,  $NR^9_2$ ,  $CO_2R^9$ ,  $CONR^9_2$ ,

N(C=NH)NH-, Het or Ar;  $R^{17} = R^{72} = H$ ,  $C_{1-6alkyl}$ ,  $R^{10}$ ,  $R^{10}$ C(O)-,  $R^{10}$ C(S)-,  $R^{10}$ OC(O)-;

```
\begin{split} R^{21} = R^{26} = & \text{C}_{5\text{-}6}\text{alkyl}; \text{C}_{2\text{-}6}\text{alkenyl}; \text{C}_{3\text{-}11}\text{cycloalkyl}; \text{T-C}_{3\text{-}6}\text{alkyl}; \text{V-C}_{1\text{-}6}\text{alkyl}; \text{T-C}_{2\text{-}6}\text{alkenyl}; \\ \text{T-} (\text{CH}_{2})_{n}\text{CH}(\text{T})(\text{CH}_{2})_{n}; \text{ optionally substituted by one or} \\ \text{two halogens, SR}^{20}, \text{OR}^{20}, \text{NR}^{20}\text{R}^{27} \text{ or C}_{1\text{-}4}\text{alkyl}; \end{split}
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5  $R^{27} = R^{28}CO, R^{28}OCO;$ 

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$$\begin{split} R^{28} = C_{1\text{-}6alkyl}; & C_{3\text{-}11} \text{cycloalkyl}; \text{ Ar; Het; T-C}_{1\text{-}6alkyl}; \\ & T\text{-}(CH_2)_n \text{CH}(T)(CH_2)_n; \text{ optionally substituted by one or} \\ & \text{two halogens, SR}^{20}, \text{ OR}^{20}, \text{ NR}^{20}\text{R}^{73}, \text{ C}_{1\text{-}6alkyl}; \end{split}$$

 $R^{20} = R^{22} = R^{23} = R^{24} = R^{25} = R^{73} = H$ ,  $C_{1-4}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

 $R^{29} =$ 

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Cbz-leucinyl-; 2-, 3-, or 4-pyridyl methyloxycarbonyl-leucinyl-; 4-imidazole acetyl-leucinyl-, phenyl acetyl-leucinyl, N,N-dimethyl-glycinyl leucinyl, 4-pyridyl acetyl-leucinyl, 2-pyridyl sulfonyl-leucinyl, 4-pyridyl carbonyl-leucinyl, acetyl-leucinyl, benzoyl-leucinyl, 4-phenoxy-benzoyl-, 2- or 3-benzyloxybenzoyl-, biphenyl acetyl, !pha- isobutyl-biphenyl acetyl, Cbz-phenylalaninyl, Cbz-norleucinyl-, Cbz-norvalinyl-, Cbz-glutamyl-, Cbz-

epsilon- (t-butyl ester)-glutamyl; acetyl-leucinyl-, 6- or 8- quinoline carbonyl, biphenyl acetyl, alpha- isobutyl-biphenyl acetyl, acetyl, benzoyl, 2- or 3- benzyloxy benzoyl, 4-phenoxy benzoyl-, Cbz-amino acid-; 2-,3-, or 4-pyridylmethyloxycarbonyl-aminoacid-; aryl C0-C6alkyloxy carbonyl-amino acid-, heteroaryl C0-C6alkyloxy carbonyl-amino acid-, aryl C0-C6alkyloxy carbonyl-amino acid-, heteroaryl C0-C6alkyloxy carbonyl-amino acid-, C1-C6alkyloxy carbonyl-amino acid-; C1-C6alkyloxy carbonyl, aryl C0-C6alkyl carbonyl, heteroaryl C0-C6alkyl carbonyl, aryl C0-C6alkyl carbonyl, heteroaryl C0-C6alkyl carbonyl, aryl C0-C6alkyl sulfonyl, aryl C0-C6alkyl sulfonyl, heteroarylC0-C6alkyl sulfonyl, aryl C0-C6alkyl sulfonyl, heteroarylC0-C6alkyl sulfonyl, aryl C0-C6alkyl sulfonyl, heteroarylC0-C6alkyl sulfonyl, aryl C0-C6alkyl sulfonyl,

 $R^{30} = -H, C_{1-6}$  alkyl;

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 $R^{31} =$ 

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Cbz-leucinyl-; 2-, 3-, or 4-pyridyl methyloxycarbonyl-leucinyl-; 4-imidazole acetyl-leucinyl-, phenyl acetyl-leucinyl, N,N-dimethyl-glycinyl leucinyl, 4-pyridyl acetyl-leucinyl, 2-pyridyl sulfonyl-leucinyl, 4-pyridyl carbonyl-leucinyl, acetyl-leucinyl, benzoyl-leucinyl, 4-phenoxy-benzoyl-, 2- or 3-benzyloxybenzoyl-, biphenyl acetyl, alpha- isobutyl-biphenyl acetyl, Cbz-phenylalaninyl, Cbz-norleucinyl-, Cbz-norvalinyl-, Cbz-glutamyl-, Cbz-

epsilon- (t-butyl ester)-glutamyl; acetyl-leucinyl-, 6- or 8- quinoline carbonyl, biphenyl acetyl, alpha- isobutyl-biphenyl acetyl, acetyl, benzoyl, 2- or 3- benzyloxy benzoyl, 4-phenoxy benzoyl-, Cbz-amino acid-; 2-,3-, or 4- pyridylmethyloxycarbonyl-aminoacid-; aryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-, aryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-, C<sub>1</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-, C<sub>1</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-; C<sub>1</sub>-C<sub>6</sub>alkyl carbonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl carbonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl carbonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl carbonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl,

 $R^{32} = OCH_2Ar, \ OCH2C_{1-6}alkyl, \ aryl \ substituted \ C_{0-6}alkyl,$  heteroaryl substituted  $C_{0-6}alkyl, 4$ -imidazole methylene; 2-, 3-, or 4-pyridylmethylneneoxy; 4-pyridyl methylene, 2-pyridyl sulfonyl, 4-pyridyl, aryl substituted  $C_{0-6}alkyloxy,$  heteroaryl substituted  $C_{0-6}alkyloxy;$ 

 $R^{33} = C_{1-6}alkyl, -CH_{2}Ph, -CH_{2}CH_{2}CO_{2}R^{34};$ 

 $R^{34} = -H, C_{1}-6alkyl;$ 

 $R^{35} = Ar, HetAr;$ 

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 $R^{36}$  = Aryl, heteroaryl, pyridyl, isoquinolinyl;

 $R^{37} = C_{1}-6alkyl$ ,  $-CH_{2}Ph$ ,  $-CH_{2}CH_{2}CO_{2}R^{34}$ ;

 $R^{38} = Cbz$ ;  $C_{1}$ -6alkyl or aryl substituted

Cbz;  $C_{1}$ -6alkyl -CO; benzcyl;  $C_{1}$ -6alkyl or aryl

substituted benzoyl;

 $R^{39} =$ 

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Cbz-leucinyl-; 2-, 3-, or 4-pyridyl methyloxycarbonyl-leucinyl-; 4-imidazole acetyl-leucinyl-, phenyl acetyl-leucinyl, N,N-dimethyl-glycinyl leucinyl, 4-pyridyl acetyl-leucinyl, 2-pyridyl sulfonyl-leucinyl, 4-pyridyl carbonyl-leucinyl, acetyl-leucinyl, benzoyl-leucinyl, 4-phenoxy-benzoyl-, 2- or 3-benzyloxybenzoyl-, biphenyl acetyl, alpha- isobutyl-biphenyl acetyl, Cbz-phenylalaninyl, Cbz-norleucinyl-, Cbz-norvalinyl-, Cbz-glutamyl-, Cbz-

epsilon- (t-butyl ester)-glutamyl; acetyl-leucinyl-, 6- or 8- quinoline carbonyl, biphenyl acetyl, alpha- isobutyl-biphenyl acetyl, acetyl, benzoyl, 2- or 3- benzyloxy benzoyl, 4-phenoxy benzoyl-, Cbz-amino acid-; 2-,3-, or 4-pyridylmethyloxycarbonyl-aminoacid-; aryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-,aryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-,heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-, C<sub>1</sub>-C<sub>6</sub>alkyloxy carbonyl-amino acid-; C<sub>1</sub>-C<sub>6</sub>alkyl carbonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl carbonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl carbonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl carbonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, heteroaryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl, aryl C<sub>0</sub>-C<sub>6</sub>alkyl sulfonyl,

 $R^{40} = H$  and  $C_{1-6}$ alkyl;

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 $R^{41} = H$  and  $C_{1-6}$ aikyl;

 $R^{42} = C_{1}$ -6alkyl, aryl substituted  $C_{1}$ -6alkyl and hetero aryl substituted  $C_{1}$ -6alkyl.; H when  $R^{43}$  is  $C_{1}$ -6alkyl, aryl substituted  $C_{1}$ -6alkyl; and heteroaryl substituted  $C_{1}$ -6alkyl;

 $R^{43} = C_{1\text{-}6alkyl}, \text{ aryl substituted } C_{1\text{-}6alkyl} \text{ and hetero aryl}$  substituted  $C_{1\text{-}6alkyl}$ ; H when  $R^{42}$  is  $C_{1\text{-}6alkyl}$ , aryl substituted  $C_{1\text{-}6alkyl}$ ; and heteroaryl substituted  $C_{1\text{-}6alkyl}$ ;

 $R^{44} = CH(R^{53})NR^{45}R^{54}$ ,  $CH(R^{55})Ar$ ,  $C_{5-6}alkyl$ ;

 $R^{45} = R^{46} = R^{47} = R^{48} = R^{49} = R^{50} = R^{51} = H$ ,  $C_{1-6}$ alkyl,  $C_{0-6}$ alkyl,  $C_{0-6}$ alkyl,  $C_{0-6}$ alkyl,  $C_{0-6}$ alkyl,

 $R^{52} = Ar$ , Het,  $CH(R^{56})Ar$ ,  $CH(R^{56})OAr$ ,  $N(R^{56})Ar$ ,  $C_{1-6}alkyl$ ,  $CH(R^{56})NR^{46}R^{57}$ ;

 $R^{53}=C_{2-6}$ alkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl,  $R^{53} \ \, \text{and} \,\, R^{45} \ \, \text{may be connected to form a pyrrolidine or piperidine ring;}$ 

 $R^{54} = R^{57} = R^{47}$ ,  $R^{47}C(O)$ ,  $R^{47}C(S)$ ,  $R^{47}OC(O)$ :

 $R^{55} = R^{56} = R^{58} = R^{59} = H, C_{1-6}alkyl, Ar-C_{0-6}alkyl,$ 

Het-C<sub>0-6</sub>alkyl;

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 $R^{60} = R^{61} = R^{62} = R^{63} = R^{64} = H, C_{1-6}$ alkyl, Ar-C<sub>0-6</sub>alkyl, or Het-C<sub>0-6</sub>alkyl;

 $R^{65} = C_{1-6}$ alkyl, Ar. Het, CH( $R^{69}$ )Ar, CH( $R^{69}$ )OAr, N( $R^{69}$ )Ar, CH( $R^{69}$ )NR61 $R^{70}$ ;

 $R^{66} = R^{69} = R^{71} = H$ ,  $C_{1-6}$ alkyl,  $(CH_2)_{0-6}$ - $C_{3-6}$ cycloalkyl, Ar- $C_{0-6}$ alkyl, Het- $C_{0-6}$ alkyl;

R<sup>67</sup> = C<sub>1-6</sub>alkyl, (CH<sub>2</sub>)<sub>0-6</sub>-C<sub>3-6</sub>cycloalkyl, Ar-C<sub>0-6</sub>alkyl,

Het-C<sub>0-6</sub>alkyl; R<sup>66</sup> and R<sup>67</sup> may be combined to form
a 3-7 membered monocyclic or 7-10-membered bicyclic
carbocyclic or heterocyclic ring, optionally substituted with
1-4 of C<sub>1-6</sub>alkyl, Ph-C<sub>0-6</sub>alkyl, Het-C<sub>0-6</sub>alkyl, C<sub>1-6</sub>alkoxy,
Ph-C<sub>0-6</sub>alkoxy, Het-C<sub>0-6</sub>alkoxy, OH, (CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>,
O(CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>:

 $R^{68} = R^{70} = R^{62}$ ,  $R^{62}C(O)$ ,  $R^{62}C(S)$ ,  $R^{62}OC(O)$ ,  $R^{62}OC(O)$ ,  $R^{62}OC(O)$ 

and pharmaceutically acceptable salts thereof.

The compounds of Formula I are hydrazidyl, bis-hydrazidyl and bisaminomethyl carbonyl compounds having in common key structural features required of protease substrates, most particularly cathepsin K substrates. These structural features endow the present compounds with the appropriate molecular shape necessary to fit into the enzymatic active site, to bind to such active site,

thereby blocking the site and inhibiting enzymatic biological activity. Referring to Formula I, such structural features include the central electrophilic carbonyl, a peptidyl or peptidomimetic molecular backbone on either side of the central carbonyl, a terminal carbobenzyloxy moiety (e.g., Cbz-leucinyl), or a mimic thereof, on the backbone on one or both sides of the carbonyl, and optionally, an isobutyl side chain extending from the backbone on one or both sides of the carbonyl.

Abbreviations and symbols commonly used in the peptide and chemical arts are used herein to describe the compounds of the present invention. In general, the amino acid abbreviations follow the IUPAC-IUB Joint Commission on Biochemical Nomenclature as described in *Eur. J. Biochem.*, 158, 9 (1984). The term "amino acid" as used herein refers to the D- or L- isomers of alanine, arginine, asparagine, aspartic acid, cysteine, glutamine, glutamic acid, glycine, histidine, isoleucine, leucine, lysine, methionine, phenylalanine, proline, serine, threonine, tryptophan, tyrosine and valine.

"C1-6alkyl" as applied herein is meant to include substituted and unsubstituted methyl, ethyl, n-propyl, isopropyl, n-butyl, isobutyl and t-butyl, pentyl, n-pentyl, isopentyl, neopentyl and hexyl and the simple aliphatic isomers thereof. Any C1-6alkyl group may be optionally substituted independently by one or two halogens, SR', OR', N(R')2, C(O)N(R')2, carbamyl or C1-4alkyl, where R' is C1-6alkyl. C0alkyl means that no alkyl group is present in the moiety. Thus, Ar-C0alkyl is equivalent to Ar.

"C3-11cycloalkyl" as applied herein is meant to include substituted and unsubstituted cyclopropane, cyclobutane, cyclopentane, cyclohexane, cyclohexane, cyclohexane, cyclohexane, cyclononane, cyclodecane, cycloundecane.

"C2-6 alkenyl" as applied herein means an alkyl group of 2 to 6 carbons wherein a carbon-carbon single bond is replaced by a carbon-carbon double bond. C2-6alkenyl includes ethylene, 1-propene, 2-propene, 1-butene, 2-butene, isobutene and the several isomeric pentenes and hexenes. Both cis and trans isomers are included.

"C2-6alkynyl" means an alkyl group of 2 to 6 carbons wherein one carboncarbon single bond is replaced by a carbon-carbon triple bond. C2-6 alkynyl includes acetylene, 1-propyne, 2-propyne, 1-butyne, 2-butyne, 3-butyne and the simple isomers of pentyne and hexyne.

"Halogen" means F, Cl, Br, and I.

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"Ar" or "aryl" means phenyl or naphthyl, optionally substituted by one or more of Ph-C<sub>0-6</sub>alkyl, Het-C<sub>0-6</sub>alkyl, C<sub>1-6</sub>alkoxy, Ph-C<sub>0-6</sub>alkoxy, Het-C<sub>0-6</sub>alkoxy, OH, (CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>, O(CH<sub>2</sub>)<sub>1-6</sub>NR<sup>58</sup>R<sup>59</sup>; where R<sup>58</sup>, R<sup>59</sup> is H, C<sub>1-6</sub>alkyl, Ar-C<sub>0-6</sub>alkyl; Het-C<sub>0-6</sub>alkyl, from C<sub>1-4</sub>alkyl, OR', N(R')<sub>2</sub>, SR', CF<sub>3</sub>, NO<sub>2</sub>, CN, CO<sub>2</sub>R', CON(R'), F, Cl, Br and I.

As used herein "Het" or "heterocyclic" represents a stable 5- to 7-membered monocyclic or a stable 7- to 10-membered bicyclic heterocyclic ring, which is either saturated or unsaturated, and which consists of carbon atoms and from one to three heteroatoms selected from the group consisting of N, O and S, and wherein the nitrogen and sulfur heteroatoms may optionally be oxidized, and the nitrogen heteroatom may optionally be quaternized, and including any bicyclic group in which any of the above-defined heterocyclic rings is fused to a benzene ring. The heterocyclic ring may be attached at any heteroatom or carbon atom which results in the creation of a stable structure, and may optionally be substituted with one or two moieties selected from C<sub>1-4</sub>alkyl, OR', N(R')<sub>2</sub>, SR', CF<sub>3</sub>, NO<sub>2</sub>, CN, CO<sub>2</sub>R', CON(R'), F, Cl, Br and I, where R' is C1-6alkyl. Examples of such heterocycles include piperidinyl, piperazinyl, 2-oxopiperazinyl, 2-oxopiperidinyl, 2oxopyrrolodinyl, 2-oxoazepinyl, azepinyl, pyrrolyl, 4-piperidonyl, pyrrolidinyl, pyrazolyl, pyrazolidinyl, imidazolyl, pyridyl, pyrazinyl, oxazolidinyl, oxazolinyl, oxazolyl, isoxazolyl, morpholinyl, thiazolidinyl, thiazolyl, quinuclidinyl, indolyl, quinolinyl, isoquinolinyl, benzimidazolyl, benzopyranyl, benzoxazolyl, furyl, pyranyl, tetrahydrofuryl, tetrahydropyranyl, thienyl, benzoxazolyl, thiamorpholinyl sulfoxide, thiamorpholinyl sulfone, and oxadiazolyl.

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"HetAr" or "heteroary!" means any heterocyclic moiety encompassed by the above definition of Het which is aromatic in character, e.g., pyridine.

It will be appreciated that the heterocyclic ring,  $\mathbb{Z}$ , includes thiazoles, oxazoles, triazoles, thiadiazoles, oxadiazoles, isoxazoles, isothiazols, imidazoles, pyrazines, pyridazines, pyrimidines, triazines and tetrazines which are available by routine chemical synthesis and are stable. The single and double bonds (i.e., --) in such heterocycles are arranged based upon the heteroatoms present so that the heterocycle is aromatic (e.g., it is a heteroary) group). The term heteroatom as applied herein refers to oxygen, nitrogen and sulfur. When the heteroaryl group comprises a five membered ring, W is preferably an electron withdrawing group, such as halogen, -CN,  $-\text{CF}_3$ ,  $-\text{NO}_2$ ,  $-\text{COR}^7$ ,  $-\text{CO}_2\text{R}^6$ ,  $-\text{CONHR}^6$ ,  $-\text{SO}_2\text{NHR}^6$ ,

NHSO<sub>2</sub>R<sup>6</sup>, -NHCOR<sup>7</sup>, -O-COR<sup>6</sup>, -SR<sup>6</sup> or NR'R<sup>6</sup>, or a similar electron withdrawing substituent as known in the art.

Certain radical groups are abbreviated herein. t-Bu refers to the tertiary butyl radical, Boc refers to the t-butyloxycarbonyl radical, Fmoc refers to the fluorenylmethoxycarbonyl radical, Ph refers to the phenyl radical, Cbz refers to the benzyloxycarbonyl radical.

Certain reagents are abbreviated herein. DCC refers to dicyclohexylcarbodiimide, DMAP is 2,6-dimethylaminopyridine, EDC refers to Nethyl-N'(dimethylaminopropyl)-carbodiimide. HOBT refers to 1-

hydroxybenzotriazole, DMF refers to dimethyl formamide, BOP refers to benzotriazol-1-yloxy-tris(dimethylamino)phosphonium hexafluorophosphate,
 DMAP is dimethylaminopyridine, Lawesson's reagent is 2,4-bis(4-methoxyphenyl)-1,3-dithia-2,4-diphosphetane-2,4-disulfide, NMM is N-methylmorpholine, TFA refers to trifluoroacetic acid, TFAA refers to trifluoroacetic anhydride and THF
 refers to tetrahydrofuran. Jones reagent is a solution of chromium trioxide, water, and sulfuric acid well-known in the art.

Compounds of formula (I) are prepared according to the methods detailed in Schemes 1-25.

#### Scheme 1

a) i-BuOCOCI, NMM, CH<sub>2</sub>N<sub>2</sub>, EtOAc, Et<sub>2</sub>O; b) HBr, AcOH, EtOAc, Et<sub>2</sub>O; c) H<sub>2</sub>NCSCO<sub>2</sub>Et, EtOH; d) NaOH, H<sub>2</sub>O, THF; e) i-BuOCOCI, NMM, NH<sub>2</sub>, THF or BOP, Et<sub>3</sub>N, RNH<sub>2</sub>, CH<sub>2</sub>Ct<sub>2</sub>; f) TFAA, pyridine, CH<sub>2</sub>Ct<sub>2</sub>; g) R<sup>4</sup>OH, Boc<sub>2</sub>O, Pyridine or R<sup>4</sup>OH, EDCI, CH<sub>2</sub>Cl<sub>2</sub>; h) piperidine, DMF; i) BOP, Et<sub>3</sub>N, D-CO<sub>2</sub>H, CH<sub>2</sub>Cl<sub>2</sub>

#### Scheme 1A

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5 a) MeI, THF; b) R'NH2, i-PrOH; c) Bromomethyl ketone, EtOH

#### Scheme 2

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a) i-BuOCOCl, NMM, NH<sub>3</sub>, THF; b) Lawesson's reagent, THF; c) BrCH<sub>2</sub>COCO<sub>2</sub>Et, TFAA, Pyridine, CH<sub>2</sub>Cl<sub>2</sub>; d) TFA; e) DCO<sub>2</sub>H, EDC•HCl, HOBT, Et<sub>3</sub>N, DMF; f) NaOH, H<sub>2</sub>O, THF

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## Scheme 2A

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a) Boc-amino acid, EDC•HCl, 1-HOBT, DMF; b) TFA; c) R<sup>5</sup>OCOCl, i-Pr<sub>2</sub>NEt

- a) Boc<sub>2</sub>O, Et<sub>3</sub>N, THF; b) hydrazine hydrate, MeOH; c) EtO<sub>2</sub>CCOCl, Pyridine.
- 5 CH<sub>2</sub>Cl<sub>2</sub>; d) Lawesson's reagent, toluene; e) TFA, CH<sub>2</sub>Cl<sub>2</sub>; f) DCO<sub>2</sub>H, EDC•HCl/HOBT, Et<sub>3</sub>N, DMF

WO 97/16177

# Scheme 4

- a) SOCl<sub>2</sub>, pyridine, Et<sub>2</sub>O, toluene; b) TFA, CH<sub>2</sub>Cl<sub>2</sub>; c) DCO<sub>2</sub>H, EDC•HCl/HOBT,
- 5 Et<sub>3</sub>N, DMF; d) NH<sub>3</sub>, EtOH

a) EDC•HCl/HOBT, Et<sub>3</sub>N, DMF; b) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, MeOH; c) CSCl<sub>2</sub>, Et<sub>3</sub>N, CHCl<sub>3</sub>

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Scheme 6

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a) H2NCS2 NH4+, EtOH; b) H2NCSNH2. EtOH

a) Et<sub>2</sub>NO; b) H<sub>2</sub>NCH<sub>2</sub>CH(NH<sub>2</sub>)CO<sub>2</sub>H

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#### Scheme 8

$$LCO_2H \xrightarrow{a} \qquad U \xrightarrow{N_2} \qquad D \xrightarrow{b} \qquad U \xrightarrow{Br} \qquad CO_2Er$$

$$1 \qquad 2 \qquad 3 \qquad 4$$

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a) i. *i*-BuOCOCI, NMM, THF; ii. CH<sub>2</sub>N<sub>2</sub>, Et<sub>2</sub>O; b) HBr, AcOH, Et<sub>2</sub>O; c) H<sub>2</sub>NCSCO<sub>2</sub>Et, EtOH; d) R<sup>63</sup>NHNH<sub>2</sub>, EtOH; e) R<sup>65</sup>CO<sub>2</sub>H, EDC•HCI, 1-HOBT, DMF.

LCO<sub>2</sub>H 
$$\xrightarrow{a}$$
 LCONH<sub>2</sub>  $\xrightarrow{b}$  LCSNH<sub>2</sub>  $\xrightarrow{c}$   $\xrightarrow{c}$   $\xrightarrow{d}$  CO<sub>2</sub>Et  $\xrightarrow{d}$ 

1 2 3 4

S

CONHNH<sub>2</sub>  $\xrightarrow{e \text{ or } f}$   $\xrightarrow{N}$   $\xrightarrow{N}$  JR<sup>55</sup>

6 (J = CO, SO2)

a) i-BuOCOCl, NMM, NH<sub>3</sub>, THF; b) Lawesson's reagent, THF; c) i. EtO<sub>2</sub>CCOCH<sub>2</sub>Br; ii. TFAA, Py, CH<sub>2</sub>Cl<sub>2</sub>; d) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, EtOH; e) R<sup>65</sup>SO<sub>2</sub>Cl, Py, CH<sub>2</sub>Cl<sub>2</sub>; f) R<sup>65</sup>CO<sub>2</sub>H, EDC•HCl, 1-HOBT, DMF.

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a) EDC•HCI, HOBT, DMF; b) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, EtOH; c) R<sup>14</sup>-B-CO<sub>2</sub>H, EDC•HCL, HOBT, DMF

a) EDC.HCl, 1-HOBT, DMF

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#### Scheme 12

a)  ${
m H_2NNH_2 \cdot H_2O}$ , MeOH; b)  ${
m Cl_2CO}$ . PhMe; c)  ${
m H_2NNH_2 \cdot H_2O}$ . MeOH; d)  ${
m R^{49}CO_2H}$ , EDC·HCI, 1-HOBT, DMF; e)  ${
m R^{52}SO_2CI}$  or  ${
m R^{52}COCI}$ , pyridine, DMF; f)  ${
m R^{52}CO_2COR^{52}}$ ; g)  ${
m R^{52}CONR^{51}NH_2}$ 

#### Scheme 12A

a) i. PhCHO, EtOH; ii. BH3·THF; b)  $\text{Cl}_2\text{CO}$ , PhMe; c)  $\text{H}_2\text{NNH}_2\cdot\text{H}_2\text{O}$ , MeOH; d)  $\text{R}^{52}\text{CO}_2\text{H}$ , EDC·HCl, 1-HOBT, DMF; e)  $\text{R}^{52}\text{SO}_2\text{Cl}$  or  $\text{R}^{52}\text{COCl}$ , pyridine, DMF; f)  $\text{R}^{52}\text{CO}_2\text{COR}^{52}$ 

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# Scheme 13

a) HBTU, NMM, DMF; b) Jones, acetone

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15 a) NMM, DMF; b) Jones, acetone

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## 5 a) EDCI, HOBT, DMF; b) NMM, DMF, 3) Jones, acetone

#### Scheme 17

a) NaN3, MeOH, H2O; b) Tosyl chloride, triethylamine, CH2Cl2; c)
 Ellman dihydropyran resin (3), PPTS, Cl(CH2)2Cl; d) PhCH2NH2, toluene,
 80 degrees C; e) HATU, N-methyl morpholine, NMP; f) HS(CH2)3SH,
 MeOH, Et3N; g) Cbz-leucine (6), HBTU, N-methyl morpholine, NMP; h)
 TFA, CH2Cl2, Me2S; i) Jones reagent, acetone

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15 a) 4-pyridyl methyl amine, isopropano!, reflux; b) Cbz-leucine, HBTU, N-methyl morpholine, DMF; c) hydrazine, MeOH, reflux; d) 2dibenzofuransulfonyl chloride, N-methyl morpholine, DMF; e) Jones reagent, acetone

#### Scheme 19

5 a) KOH, MeOH/H2O; b) R<sup>66</sup>NHNH<sub>2</sub>, EtOH; c) EDC•HCl, 1-HOBT, DMF

#### Scheme 20

$$EtO_{2}CCOCH_{2}Br \xrightarrow{a} H_{2}N \xrightarrow{N} CO_{2}Et \xrightarrow{b} Br \xrightarrow{N} CO_{2}Et$$

$$1 \qquad 2 \qquad 3$$

$$Cord Ar \xrightarrow{N} CO_{2}Et \xrightarrow{e} Ar \xrightarrow{N} CONHNH_{2} \xrightarrow{f}$$

$$4 \qquad 5$$

$$Ar \xrightarrow{N} N \xrightarrow{N} N \xrightarrow{N} R^{85}$$

$$6$$

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a) Thiourea, EtOH; b) i. NaNO<sub>2</sub>, 16% aqueous HBr; ii. CuBr, 16% aqueous HBr; iii. HBr (cat.), EtOH; c) ArB(OH)<sub>2</sub>, Pd(PPh<sub>3</sub>)<sub>4</sub>, CsF, DME; d) ArSnMe<sub>3</sub>, Pd(PPh<sub>3</sub>)<sub>4</sub>, PhMe; e) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, EtOH; e) R<sup>65</sup>CO<sub>2</sub>H, EDC•HCl, 1-HOBT, DMF.

#### Scheme 21

a) R<sup>67</sup>NH<sub>2</sub>, Py, CH<sub>2</sub>Cl<sub>2</sub>; b) LiAlH<sub>4</sub>, THF; c) i. Cl<sub>2</sub>CS, Py, CH<sub>2</sub>Cl<sub>2</sub>; ii. NH<sub>3</sub>, MeOH or I. PhCONCS, CHCl<sub>3</sub>; ii. K<sub>2</sub>CO<sub>3</sub>, MeOH, H<sub>2</sub>O; d) EtO<sub>2</sub>CCOCH<sub>2</sub>Br, EtOH; e) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, EtOH; e) R<sup>65</sup>CO<sub>2</sub>H, EDC•HCl, 1-HOBT, DMF.

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Scheme 22

a) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, EtOH; 5) LCO<sub>2</sub>CO<sub>2</sub>i-Bu, 200 °C; c) H<sub>2</sub>NNH<sub>2</sub>•H<sub>2</sub>O, EtOH; d) R<sup>65</sup>CO<sub>2</sub>H, EDC•HCl, 1-HOBT, DMF

#### Scheme 23

a) TFA; b) R<sup>62</sup>CO<sub>2</sub>H, EDC•HCl, 1-HOBT, DMF; c) R<sup>62</sup>SO<sub>2</sub>Cl, *i*-Pr<sub>2</sub>NEt

#### Scheme 24

a) EDCI, DMF; b) 2-PhCH<sub>2</sub>OPhSO<sub>2</sub>Cl, NMM, DMF; c) TFA, DCM; d) 4-pyridyl acetic acid, HBTU, NMM, DMF; e) Jones

#### Scheme 25

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a) HBTU, NMM, DMF, allyl amine; b) mCPBA, DCM; c) MeNH<sub>2</sub>, isopropanol, 70 C; d) Cbz-leucine, EDCI, DMF; e) Jones, acetone

In another aspect, the present invention provides a novel cysteine protease in crystalline form, as defined by the positions in Table I herein.

In still another aspect, the present invention provides a novel protease composition characterized by a three dimensional catalytic site formed by the atoms of the amino acid residues listed in Table XXIX herein.

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The three dimensional (3D) structure of the instant protease reveals that human cathepsin K is highly homologous to other known cysteine proteinases of the papain family. Cathepsin-K folds into two subdomains separated by the active site cleft, a characteristic of the papain family of cysteine proteases. The overall fold of cathepsin K is very similar to that of papain and actinidin. There is an insertion of one additional residue in cathepsin K at residue alanine 79 compared to papain. This insertion is easily accommodated in the turn at the carboxy terminal end of the helix formed by residues methionine 68-lysine 77 of cathepsin K. There is a different conformation for the backbone atoms of residues asparagine 99 to lysine 103 at the surface of cathepsin K compared to that in papain. Other differences in the backbone conformations between cathepsin K and papain are: a two residue insertion in loop residues 126-127, a two residue insertion at residue aspartate 152, the insertion of 4 residues at glutamine 172 and a difference in the conformation of the loop around residue lysine 200. There are many more differences in the structure of human cathepsin K and human cathepsin B, however, the secondary structure is preserved well between these two enzymes.

Listed in Figure 1 are the known amino acid sequences for the papain superfamily of cysteine proteases cathepsin K, cathepsin S, cathepsin L, papain, actinidin, cathepsin H and cathepsin B, aligned to illustrate the homologies there between.

According to the present invention the crystal structure of human cathepsin K has been determined in the absence of inhibitor and in complex with nine separate inhibitors at resolutions from 3.0 to 2.2 Ångstroms. The structures were determined using the method of molecular replacement and refined to R<sub>c</sub> values ranging from 0.190-0.267 with the exception of the enzyme in the absence of inhibitor which was not refined.

Further refinement of the atomic coordinates will change the numbers in Table I. Refinement of the crystal structure from another crystal form will result in a new set of coordinates, determination of the crystal structure of another cysteine

protease will also result in different set of numbers for coordinates in Table I which has an experimental error of approximately 0.4 Ångstroms. Also for example, the amino acid sequence of the cysteine proteases can be varied by mutation derivatization or by use of a different source of the protein.

Human cathepsin K contains 215 amino acids and the model of the enzyme provided herein is represented by all 215 residues.

The cathepsin K crystal structure reveals an active site that is heretofor unknown and comprises a distinct three dimensional arrangement of atoms.

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Table I discloses the protein coordinates of cathepsin K. These data are reported for the crystal structures described herein. The data are reported in Ångstroms with reference to an orthogonal coordinate system in standard format, illustrating the atom, i.e., nitrogen, oxygen, carbon, sulfur (at  $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\delta$ , or  $\epsilon$ , positions in the amino acid residues); the amino acid residue in which the atom is located with amino acid number, and the coordinates X, Y and Z in Angstroms (Å) from the crystal structure. Note that each atom in the active site and the entire structure has an unique position in the crystal. The data also report the B or Temperature Factor values, which indicate the degree of thermal motion of the atom in root mean square displacement measurements (Å<sup>2</sup>). Figure 2 illustrates the cathepsin K structure of the invention, including the active site.

The active site of cathepsin K bound to E-64 is shown in Figure 3. The conformation of E-64 bound to cathepsin K resembles that seen in the published structures of the papain-E-64 complex (Varughese, K.I., Biochemistry 28, 1330-1332 (1989)) and actinidin-E-64 Varughese, K.I., Biochemistry 31, 5172-5176 (1992)). The covalent bond between the sulfur of cysteine 25 and the carbon C2 of the inhibitor is very clear in the electron density. Differences in the sidechain atoms lining the active site pockets on the enzyme of the various members of the papain family of cysteine proteases give rise to different interactions between the atoms of E-64 and the protein in these structures. In cathepsin K, the isobutyl atoms of the leucine lie well buried in the hydrophobic pocket formed by the side chain atoms of the cathepsin K residues leucine 160, alanine 134 and methionine 68 shielding these atoms of E-64 from solvent. In papain the leucyl side chain atoms of E-64 do not penetrate as deeply into this hydrophobic pocket. Another pocket of cathepsin K is occupied by the guanidinium atoms of E-64. A hydrogen bond forms between N4 of E-64 and the backbone carbonyl oxygen of glutamate 59 and the OD2 oxygen of aspartate 61. The carboxylate oxygen of aspartate 61 also makes a hydrogen bond

with the N3 atom of E-64. The sidechain atoms of aspartate 61 lie at the entrance to this pocket in cathepsin K. These interactions are not possible in papain because the corresponding residue in papain is tyrosine 61 which blocks access. The carboxylate oxygens of E-64 make hydrogen bonding interactions with the ND1 atom of histidine 162 and the NE2 atom of glutamine 19. These interactions are also seen in 5 papain and actinidin. The atoms of E-64 do not penetrate the complete region of the enzyme active site. As in papain, the backbone nitrogen atoms of residue glycine 66 in cathepsin K makes a hydrogen bond with the carbonyl oxygen atom O4 of the E-64. Also, the carbonyl oxygen of glycine 66 of cathepsin K forms a hydrogen bond with N2 of E-64. A portion of the regions of the active site are very similar in 10 conformation in cathepsin K, papain and actinindin. A comparison of the active site of cathepsin K and cathepsin B reveals many more differences than observed in comparing papain or actinidin to cathepsin K. A portion of the active site of cathepsin B differs significantly from the corresponding portion of the active site in cathepsin K. The presence of the loop glutamate 107 - proline 116 in human 15 cathepsin B is presumed responsible for the dipeptidyl carboxypeptidase activity of this enzyme and has no equivalent in cathepsin K, papain or actinidin. This loop makes this region of the active site of cathepsin B much smaller than in the other members of this papain family of cysteine proteases including cathepsin K. Despite the differences between the active sites of human cathepsin B and cathepsin K, the 20 active site cysteine residues are almost exactly superimposed by an alignment of structurally homologous alpha carbon atoms in cathepsin B and cathepsin K. Differences in the hydrophobic pocket near leucine 160 in cathepsin K are also evident in cathepsin B. The residues forming this pocket are replaced by proline 78 in place of methionine 68 in cathepsin K and glutamate 243 in cathepsin B is 25 structurally equivalent to leucine 160 in cathepsin K. Interestingly, the residues whose sidechain atoms form hydrogen bonds to the E-64 inhibitor in cathepsin K, namely histidine 162, glutamine 19 and aspartate 61, have structurally homologous residues in cathepsin B, namely histidine 197, glutamine 23 and aspartate 67 30 respectively.

Specific interactions of certain inhibitors of the present invention at the active site of cathepsin K are detailed hereinbelow.

3 (S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone makes hydrophobic contacts with the enzyme residues indole ring of tryptophan 184 and the sidechain atom CG of glutamine 19. Oxygen O26 forms a

bifurcated hydrogen bond with the amide nitrogen of cysteine 25 and the NE2 atom of glutamine 19. The active site nucleophilic sulfur of residue cysteine 25 is covalently linked to carbon C25 of the inhibitor, which adopts a tetrahedral conformation.

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Bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one exhibits the same interaction as 3 (S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone; carbon C21 of this inhibitor is covalently linked to SG of cysteine 25. The isopropyl atoms CC34,C35,C36 and C37 of the inhibitor form hydrophobic interactions with the sidechain atoms of residues on the enzyme surface, which form a hydrophobic pocket. This pocket is formed by atoms from methionine 68, leucine 209, alanine 163 and alanine 134 and portions of tyrosine 67.

2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide has interactions similar to bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one and, in addition, the atoms C23-29 of the inhibitor CBZ group make an edge-face stacking interaction with the phenol ring of tyrosine 67. Inhibitor atom C21 is covalently bound the enzyme.

The sulfur atom of (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide contacts the ND1 atom of histidine 163 and the indole ring of tryptophan 184. Carbon C22 is covalently attached to SG of cysteine 25.

The CBZ atoms C20-26 of 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide interact with the sidechain atoms of leucine 160. Carbon C19 is covalently attached to SG of cysteine 25.

Cathepsin K binds selectively one stereoisomer of 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone. Carbon C22 is covalently attached to SG of cysteine 25. Atoms C14 and C15 of the inhibitor 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone form hydrophobic contacts with the sidechain atoms of glutamine 143 and asparagine 161 and the mainchain of alanine 137 and serine 138.

4-[N-[(4-pyridylmethoxy)carbonyi]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone interacts in a similar manner to 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone. Again one stereoisomer is bound. Carbon C17 is covalently attached to SG of cysteine 25. The interaction of 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl]-3-pyrrolidinone is

the same as for 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone, except carbon C22 is covalently attached to SG of cysteine 25.

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Atom O24 of 1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxyphenyl-sulfonyl)-amino-propan-2-one forms a hydrogen bond interaction with the amide NH of glycine 66. Carbon C19 is covalently attached to SG of cysteine 25.

In summary, all inhibitors exhibit an aromatic interaction with atoms of the indole of Tryptophan 184. Isopropyl atoms C12-15 of 2,2'-N,N'-bisbenzyloxycarbonyl-L-leucinylcarbohydrazide and (1S)-N-[2-[(1-

- benzyloxy carbonylamino) 3-methylbutyl] thiazol-4-ylcarbonyl] N'-(N-methylbutyl) (N-methylbutyl) N'-(N-methylbutyl) N'-(N-m10 benzyloxycarbonyl-L-leucinyl)hydrazide make hydrophobic contacts with main chain atoms of residues glutamine 21, cysteine 22 and glycine 23. The NE2 atom of glutamine 19 is able to donate a hydrogen bond to oxygen atom 2,2'-N,N'-bisbenzyloxycarbonyl-L-leucinylcarbohydrazide:O22, 1-N-(N-imidazole acetyl-
- leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one:O20, 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-Lleucinyl)]carbohydrazide:O20, 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone:O23, bis-(Cbz-leucinyl)-1,3diamino-propan-2-one:O22, 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-
- methyl-1-(1-propoxy)-2-hexanone:O26, 4-[N-[(4-pyridylmethoxy)carbonyl]-L-20 leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone:O42, (1S, 2'R)-N-2-[[(1-benzyloxycarbonyl)amino]-3-methylbutyl]thiazol-4-ylcarbonyl-N'-2'-(benzyloxycarbonyl)amino-4'-methylpenanoylhydrazide:O23, 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-
- pyrrolidinone:O23. The backbone amide nitrogen of glycine 66 donates a hydrogen 25 bond to 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide:O39, 1-N-(Nimidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan- $\hbox{$2$-one:}O24, \hbox{$2$-[N-(3-benzyloxybenzoyl)]-$2'-[N'-(N-benzyloxycarbonyl-L-benzyloxybenzoyl)]-$2'-[N'-(N-benzyloxybenzoyl$ leucinyl)]carbohydrazide:O37, 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-
- [(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone:O40, bis-(Cbz-leucinyl)-1,3-30 diamino-propan-2-one:O39, (1S, 2'R)-N-2-[[(1-benzyloxycarbonyl)amino]-3methylbutyl]thiazol-4-ylcarbonyl-N'-2'-(benzyloxycarbonyl)amino-4'methylpenanoylhydrazide:O40, 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone:O31. The hydrophobic pocket lined with atoms from residues methionine 68, leucine 209, alanine 163 and alanine 134 and portions

of tyrosine 67 interact with the isopropyl atoms; bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one:C34-37, 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide: C34-37, (1S)-N-{2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide; :C35-38, 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide:C32-35, 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone:C35-38, 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone:C19-22, 1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one:C26-29. All inhibitors except 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone and 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl])-3-pyrrolidinone have aromatic groups that interact with tyrosine 67 on the protein. All inhibitors are covalently linked to the cysteine 25 SG atom through an inhibitor carbon atom.

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The crystal structure of the protease of the present invention reveals the three dimensional structure of novel active site formed by the atoms of the amino acid residues listed in Table XXIX.

This structure is clearly useful in the structure-based design of protease inhibitors, which may be used as therapeutic agents against diseases in which inhibition of bone resorption is indicated. The discovery of the novel cathepsin K catalytic site permits the design of potent, highly selective protease inhibitors.

Another aspect of this invention involves a method for identifying inhibitors of cathepsin K characterized by the crystal structure and novel active site described herein, and the inhibitors themselves. The novel protease crystal structure of the invention permits the identification of inhibitors of protease activity. Such inhibitors may bind to all or a portion of the active site of cathepsin K; or even be competitive or non-competitive inhibitors. Once identified and screened for biological activity, these inhibitors may be used therapeutically or prophylactically to block protease activity.

One design approach is to probe the cathepsin K of the invention with molecules composed of a variety of different chemical entities to determine optimal sites for interaction between candidate cathepsin K inhibitors and the enzyme. For example, high resolution X-ray diffraction data collected from crystals saturated with solvent allows the determination of where each type of solvent molecule sticks.

Small molecules that bind tightly to those sites can then be designed and synthesized and tested for their cathepsin K inhibitor activity.

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This invention also enables the development of compounds that can isomerize to short-lived reaction intermediates in the chemical reaction of a substrate or other compound that binds to or with cathepsin K. Thus, the time-dependent analysis of structural changes in cathepsin K during its interaction with other molecules is permitted. The reaction intermediates of cathepsin K can also be deduced from the reaction product in co-complex with cathepsin K. Such information is useful to design improved analogues of known cysteine protease inhibitors or to design novel classes of inhibitors based on the reaction intermediates of the cathepsin K enzyme and cathepsin K inhibitor co-complex. This provides a novel route for designing cathepsin K inhibitors with both high specificity and stability.

Another approach made possible by this invention, is to screen computationally small molecule data bases for chemical entities or compounds that can bind in whole, or in part, to the cathepsin K enzyme. In this screening, the quality of fit of such entities or compounds to the binding site may be judged either by shape complementarity [R. L. DesJarlais et al., <u>J. Med. Chem.</u> 31:722-729 (1988)] or by estimated interaction energy [E. C. Meng et al, <u>J. Comp. Chem.</u>, 13:505-524 (1992)].

Because cathepsin K may crystallize in more than one crystal form, the structure coordinates of cathepsin K, or portions thereof, as provided by this invention are particularly useful to solve the structure of those other crystal forms of cathepsin K. They may also be used to solve the structure of cathepsin K mutants, cathepsin K co-complexes, or of the crystalline form of any other protein with significant amino acid sequence homology to any functional domain of cathepsin K.

One method that may be employed for this purpose is molecular replacement. In this method, the unknown crystal structure, whether it is another crystal form of cathepsin K, a cathepsin K mutant, or a cathepsin K co-complex, or the crystal of some other protein with significant amino acid sequence homology to any functional domain of cathepsin K, may be determined using the cathepsin K structure coordinates of this invention as provided in Table I. This method will provide an accurate structural form for the unknown crystal more quickly and efficiently than attempting to determine such information ab initio.

Thus, the cathepsin K structure provided herein permits the screening of known molecules and/or the designing of new molecules which bind to the protease structure, particularly at the active site, via the use of computerized evaluation systems. For example, computer modeling systems are available in which the sequence of the protease, and the protease structure (i.e., atomic coordinates of cathepsin K and/or the atomic coordinate of the active site cavity, bond angles, dihedral angles, distances between atoms in the active site region, etc. as provided by Table I may be input. Thus, a machine readable medium may be encoded with data representing the coordinates of Table I in this process. The computer then generates structural details of the site into which a test compound should bind, thereby enabling the determination of the complementary structural details of said test compound.

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More particularly, the design of compounds that bind to or inhibit cathepsin K according to this invention generally involves consideration of two factors. First, the compound must be capable of physically and structurally associating with cathepsin K. Non-covalent molecular interactions important in the association of cathepsin K with its substrate include hydrogen bonding, van der Waals and hydrophobic interactions.

Second, the compound must be able to assume a conformation that allows it to associate with cathepsin K. Although certain portions of the compound will not directly participate in this association with cathepsin K, those portions may still influence the overall conformation of the molecule. This, in turn, may have a significant impact on potency. Such conformational requirements include the overall three-dimensional structure and orientation of the chemical entity or compound in relation to all or a portion of the binding site, e.g., active site or accessory binding site of cathepsin K, or the spacing between functional groups of a compound comprising several chemical entities that directly interact with cathepsin K.

The potential inhibitory or binding effect of a chemical compound with cathepsin K may be estimated prior to its actual synthesis and testing by the use of computer modeling techniques. If the theoretical structure of the given compound suggests insufficient interaction and association between it and cathepsin K, synthesis and testing of the compound is obviated. However, if computer modeling indicates a strong interaction, the molecule may then be synthesized and tested for

its ability to bind to cathepsin K in a suitable assay. In this manuer, synthesis of inoperative compounds may be avoided.

An inhibitory or other binding compound of cathepsin K may be computationally evaluated and designed by means of a series of steps in which chemical entities or fragments are screened and selected for their ability to associate with the individual binding pockets or other areas of cathepsin K.

One skilled in the art may use one of several methods to screen chemical entities or fragments for their ability to associate with cathepsin K and more particularly with the individual binding pockets of the cathepsin K active site or accessory binding site. This process may begin by visual inspection of, for example, the active site on the computer screen based on the cathepsin K coordinates in Table I. Selected fragments or chemical entities may then be position cathepsin K. Docking may be accomplished using software such as Quanta and Sybyl, followed by energy minimization and molecular dynamics with standard molecular mechanics forcefields, such as CHARMM and AMBER.

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Specialized computer programs may also assist in the process of selecting fragments or chemical entities. These include:

- GRID [P. J. Goodford, "A Computational Procedure for Determining Energetically Favorable Binding Sites on Biologically Important Macromolecules", J. Med. Chem., 28:849-857 (1985)]. GRID is available from Oxford University, Oxford, UK.
- MCSS [A. Miranker and M. Karplus, "Functionality Maps of Binding Sites: A Multiple Copy Simultaneous Search Method", Proteins: Structure, Function and Genetics, 11:29-34 (1991)]. MCSS is available from Molecular Simulations, Burlington, MA.
- AUTODOCK [D. S. Goodsell and A. J. Olsen, "Automated Docking of Substrates to Proteins by Simulated Annealing", <u>Proteins: Structure. Function</u>, and Genetics, 8:195-202 (1990)]. AUTODOCK is available from Scripps Research Institute, La Jolla, CA.
- DOCK [I. D. Kuntz et al, "A Geometric Approach to Macromolecule-Ligand Interactions", <u>J. Mol. Biol.</u>, 161:269-288 (1982)]. DOCK is available from University of California, San Francisco, CA.

Additional commercially available computer databases for small molecular compounds includes Cambridge Structural Database and Fine Chemical Database, for a review see Rusinko, A., Chem. Des. Auto. News 8, 44-47 (1993).

Once suitable chemical entities or fragments have been selected, they can be assembled into a single compound or inhibitor. Assembly may be proceeded by visual inspection of the relationship of the fragments to each other on the three-dimensional image displayed on a computer screen in relation to the structure coordinates of cathepsin K. This would be followed by manual model building using software such as Quanta or Sybyl.

Useful programs to aid one of skill in the art in connecting the individual chemical entities or fragments include:

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- CAVEAT [P. A. Bartlett et al, "CAVEAT: A Program to Facilitate the Structure-Derived Design of Biologically Active Molecules", in Molecular Recognition in Chemical and Biological Problems", Special Pub., Royal Chem. Soc. 78, pp. 182-196 (1989)]. CAVEAT is available from the University of California, Berkeley, CA.
- 3D Database systems such as MACCS-3D (MDL Information Systems, San Leandro, CA). This area is reviewed in Y. C. Martin, "3D Database Searching in Drug Design", <u>J. Med. Chem.</u>, <u>35</u>:2145-2154 (1992).
  - HOOK (available from Molecular Simulations, Burlington, MA).
    Instead of proceeding to build a cathepsin K inhibitor in a step-wise fashion one fragment or chemical entity at a time as described above, inhibitory or other type of binding compounds may be designed as a whole or "de novo" using either an empty active site or optionally including some portion(s) of a known inhibitor(s).
    These methods include:
  - LUDI [H.-J. Bohm, "The Computer Program LUDI: A New Method for the De Novo Design of Enzyme Inhibitors", J. Comp. Aid. Molec. Design, 6:61-78 (1992)]. LUDI is available from Biosym Technologies, San Diego, CA.
  - LEGEND [Y. Nishibata and A. Itai, <u>Tetrahedron</u>, <u>47</u>:8985 (1991)]. LEGEND is available from Molecular Simulations, Burlington, MA.
  - LeapFrog (available from Tripos Associates, St. Louis, MO).

Other molecular modeling techniques may also be employed in accordance with this invention. See, e.g., N. C. Cohen et al, "Molecular Modeling Software and Methods for Medicinal Chemistry", J. Med. Chem., 33:883-894 (1990). See also, M. A. Navia and M. A. Murcko, "The Use of Structural Information in Drug Design", Current Opinions in Structural Biology, 2:202-210 (1992). For example, where the structures of test compounds are known, a model of the test compound may be superimposed over the model of the structure of the invention. Numerous

methods and techniques are known in the art for performing this step, any of which may be used. See, e.g., P.S. Farmer, Drug Design, Ariens, E.J., ed., Vol. 10, pp 119-143 (Academic Press, New York, 1980); U.S. Patent No. 5,331,573; U.S. Patent No. 5,500,807; C. Verlinde, <u>Structure</u>, 2:577-587 (1994); and I. D. Kuntz, <u>Science</u>, 257:1078-1082 (1992). The model building techniques and computer evaluation systems described herein are not a limitation on the present invention.

Thus, using these computer evaluation systems, a large number of compounds may be quickly and easily examined and expensive and lengthy biochemical testing avoided. Moreover, the need for actual synthesis of many compounds is effectively eliminated.

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Once identified by the modeling techniques, the protease inhibitor may be tested for bioactivity using standard techniques. For example, structure of the invention may be used in binding assays using conventional formats to screen inhibitors. Suitable assays for use herein include, but are not limited to, the enzymelinked immunosorbent assay (ELISA), or a fluoresence quench assay. See, for example, the cathepsin K activity assay of Example 2 below. Other assay formats may be used; these assay formats are not a limitation on the present invention.

In another aspect, the protease structure of the invention permit the design and identification of synthetic compounds and/or other molecules which have a shape complimentary to the conformation of the protease active site of the invention. Using known computer systems, the coordinates of the protease structure of the invention may be provided in machine readable form, the test compounds designed and/or screened and their conformations superimposed on the structure of the protease of the invention. Subsequently, suitable candidates identified as above may be screened for the desired protease inhibitory bioactivity, stability, and the like.

Once identified and screened for biological activity, these inhibitors may be used therapeutically or prophylactically to block cathepsin K activity.

The following examples illustrate various aspects of this invention. These examples do not limit the scope of this invention which is defined by the appended claims.

#### EXAMPLE 1: Analysis of the Structure of Cathepsin K

A. Expression, Purification and Crystallization

Cathepsin K (see Fig. 1) was expressed and purified as described in

Bossard, M. J., et al., J. Biol. Chem. 271, 12517-12524 (1996).

Crystals of cathepsin K were grown by vapor diffusion in hanging drops from a solution of 30% PEG 8000, 0.1 M Na<sup>+</sup>/K<sup>+</sup> phosphate at pH 4.5 containing 0.2M Li<sub>2</sub>SO<sub>4</sub>. Crystals of the complex are tetragonal, space group P4<sub>3</sub>2<sub>1</sub>2, with cell constants of a=57.7 Ångstroms and c=131.1 Ångstroms. The crystals contain one molecule in the asymmetric unit and contain 36 % solvent with a V<sub>m</sub> value of 2.3 Å<sup>3</sup>/Dalton. The structure was determined by molecular replacement using X-PLOR [Brunger, A.T., et al., *Science*, 235, 458-460 (1987)]. The starting model consisted of the protein atoms from the cathepsin K E-64 complex structure described herein.

#### B. Model Building and Refinement

Using the three-dimensional electron density map obtained from above, the polypeptide chain of the cathepsin K can be traced without ambiguity. All 215 residues with side chains were built using the 3-D computer graphics program FRODO [Jones, T.A., J. Appl. Crystallogr., 11, 268-272 (1978)]. Each of these 215 amino acids residues was manually positioned in its electron density, allowing for a unique position for each atom in cathepsin K in which each position is defined by a unique set of atomic coordinates (X,Y,Z) as shown in Table I. Starting with these atomic coordinates, a diffraction pattern was calculated and compared to the experimental data. The difference between the calculated and experimentally determined diffraction patterns was monitored by the value of R<sub>C</sub>. The refinement (using X-PLOR) of the structural model necessitates adjustments of atomic positions to minimize the R-factor, where a value of below 20% is typical for a good quality protein structure and a value of higher than 25% usually indicates the need of further refinement.

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#### **EXAMPLE 2: Assays**

#### Determination of cathepsin K proteolytic catalytic activity

All assays for cathepsin K were carried out with human recombinant enzyme. Standard assay conditions for the determination of kinetic constants used a fluorogenic peptide substrate, typically Cbz-Phe-Arg-AMC, and were determined in 100 mM Na acetate at pH 5.5 containing 20 mM cysteine and 5 mM EDTA. Stock substrate solutions were prepared at concentrations of 10 or 20 mM in DMSO with 20 uM final substrate concentration in the assays. All assays contained 10% DMSO. Independent experiments found that this level of DMSO had no effect on enzyme activity or kinetic constants. All assays were conducted at ambient temperature.

Product fluorescence (excitation at 360 nM; emission at 460 nM) was monitored with a Perceptive Biosystems Cytofluor II fluorescent plate reader. Product progress curves were generated over 20 to 30 minutes following formation of AMC product.

#### 5 Inhibition studies

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Potential inhibitors were evaluated using the progress curve method. Assays were carried out in the presence of variable concentrations of test compound. Reactions were initiated by addition of enzyme to buffered solutions of inhibitor and substrate. Data analysis was conducted according to one of two procedures depending on the appearance of the progress curves in the presence of inhibitors. For those compounds whose progress curves were linear, apparent inhibition constants  $(K_{i,app})$  were calculated according to equation 1 (Brandt *et al.*, *Biochemistry*, 1989, 28, 140):

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$$v = V_m A / [K_a(1 + I/K_{i, app}) + A]$$
 (1)

where  $\nu$  is the velocity of the reaction with maximal velocity  $V_m$ , A is the concentration of substrate with Michaelis constant of  $K_a$ , and I is the concentration of inhibitor.

For those compounds whose progress curves showed downward curvature characteristic of time-dependent inhibition, the data from individual sets was analyzed to give  $k_{obs}$  according to equation 2:

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$$[AMC] = v_{SS} t + (v_0 - v_{SS}) [1 - exp(-k_{obs}t)] / k_{obs}$$
(2)

where [AMC] is the concentration of product formed over time t,  $v_0$  is the initial reaction velocity and  $v_{SS}$  is the final steady state rate. Values for  $k_{ObS}$  were then analyzed as a linear function of inhibitor concentration to generate an apparent second order rate constant ( $k_{ObS}$  / inhibitor concentration or  $k_{ObS}$  / [I]) describing the time-dependent inhibition. A complete discussion of this kinetic treatment has been fully described (Morrison et al., Adv. Enzymol. Relat. Areas Mol. Biol., 1988, 61, 201).

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This assay measures the affinity of inhibitors to cathepsin K. One skilled in the art would consider any compound exhibiting a  $K_i$  value of less than 50 micromolar to be a potential lead compound for further research. Preferably, the compounds used in the method of the present invention have a  $K_i$  value of less than 1 micromolar. Most preferably, said compounds have a  $K_i$  value of less than 100 nanomolar.

#### **Human Osteoclast Resorption Assay**

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Aliquots of osteoclastoma-derived cell suspensions were removed from liquid nitrogen storage, warmed rapidly at 37°C and washed x1 in RPMI-1640 medium by centrifugation (1000 rpm, 5 min at 4°C). The medium was aspirated and replaced with murine anti-HLA-DR antibody, diluted 1:3 in RPMI-1640 medium, and incubated for 30 min on ice The cell suspension was mixed frequently.

The cells were washed x2 with cold RPMI-1640 by centrifugation (1000 rpm, 5 min at 4°C) and then transferred to a sterile 15 mL centrifuge tube. The number of mononuclear cells were enumerated in an improved Neubauer counting chamber.

Sufficient magnetic beads (5 / mononuclear cell), coated with goat anti-mouse IgG, were removed from their stock bottle and placed into 5 mL of fresh medium (this washes away the toxic azide preservative). The medium was removed by immobilizing the beads on a magnet and is replaced with fresh medium.

The beads were mixed with the cells and the suspension was incubated for 30 min on ice. The suspension was mixed frequently. The bead-coated cells were immobilized on a magnet and the remaining cells (osteoclast-rich fraction) were decanted into a sterile 50 mL centrifuge tube. Fresh medium was added to the bead-coated cells to dislodge any trapped osteoclasts. This wash process was repeated x10. The bead-coated cells were discarded.

The osteoclasts were enumerated in a counting chamber, using a large-bore disposable plastic Pasteur pipette to charge the chamber with the sample. The cells were pelleted by centrifugation and the density of osteoclasts adjusted to 1.5x10<sup>4</sup>/mL in EMEM medium, supplemented with 10% fetal calf serum and 1.7g/liter of sodium bicarbonate. 3 mL aliquots of the cell suspension (per treatment) were decanted into 15 mL centrifuge tubes. These cells were pelleted by centrifugation. To each tube 3 mL of the appropriate treatment was added (diluted to 50 uM in the EMEM medium). Also included were appropriate vehicle controls, a

positive control (87MEM1 diluted to 100 ug/mL) and an isotype control (IgG2a diluted to 100 ug/mL). The tubes were incubate at 37°C for 30 min.

0.5 mL aliquots of the cells were seeded onto sterile dentine slices in a 48-well plate and incubated at 37°C for 2 h. Each treatment was screened in quadruplicate. The slices were washed in six changes of warm PBS (10 mL / well in a 6-well plate) and then placed into fresh treatment or control and incubated at 37°C for 48 h. The slices were then washed in phosphate buffered saline and fixed in 2% glutaraldehyde (in 0.2M sodium cacodylate) for 5 min., following which they were washed in water and incubated in buffer for 5 min at 37°C. The slices were then washed in cold water and incubated in cold acetate buffer / fast red garnet for 5 min at 4°C. Excess buffer was aspirated, and the slices were air dried following a wash in water.

The TRAP positive osteoclasts were enumerated by bright-field microscopy and were then removed from the surface of the dentine by sonication. Pit volumes were determined using the Nikon/Lasertec ILM21W confocal microscope.

#### **EXAMPLE 3: Method of Detecting Inhibitors**

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The three dimensional atomic structure can be readily used as a template for selecting potent inhibitors. Various computer programs and databases are available for the purpose. A good inhibitor should at least have excellent steric and electrostatic complementarity to the target, a fair amount of hydrophobic surface buried and sufficient conformational rigidity to minimize entropy loss upon binding. The approach usually comprises several steps:

- 1) Define a region to target. the active site cavity of cathepsin K can be selected, but any place that is essential to the protease activity could become a potential target. Since the crystal structure has been determined, the spatial and chemical properties of the target region is known.
- 2) Docking a small molecule onto the target. Many methods can be used to archive this. Computer databases of three-dimensional structures are available for screening millions of small molecular compounds. A negative image of these compounds can be calculated and used to match the shape of the target cavity. The profiles of hydrogen bond donor-acceptor and lipophilic points of these compounds can also be used to complement those of the target. Anyone skilled in the art would be able to identify many small molecules or fragments as hits.

3) Linking and extending recognition fragments. Using the hits identified by above procedure, one can incorporate different functional groups or small molecules into a single, larger molecule. The resulting molecule is likely to be more potent and have higher specificity. It is also possible to try to improve the "seed" inhibitor by adding more atoms or fragments that will interact with the target protein. The originally defined target region can be readily expanded to allow further necessary extension.

A limited number of promising compounds can be selected through the process. They can then be synthesized and assayed for their inhibitory properties. The success rate can sometimes be as high as 20%, and it may still be higher with the rapid progresses in computing methods.

#### **EXAMPLE 4: Crystallization of Enzyme with Inhibitors**

#### 15 A. <u>Preparation of Inhibitors</u>

Compound 1. Preparation of 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

a) 3-hydroxy-4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-pyrrolidinecarboxylic acid 1,1dimethylethyl ester

To a solution of 3-hydroxy-4-amino-1-pyrrolidinecarboxylic acid, 1,1-dimethylethyl ester (202 mg, 1.14 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (5 mL) was added CBZ-leucine (302.9 mg, 1.14 mmol), HOBT (154 mg, 1.14 mmol) and EDC (262.2 mg, 1.37 mmol). The reaction was allowed to stir until complete by TLC analysis whereupon it was diluted with EtOAc and washed sequentially with pH 4 buffer, sat. K<sub>2</sub>CO<sub>3</sub>, water and brine. The organic layer was dried (MgSO<sub>4</sub>), filtered and concentrated. Column chromatography of the residue (3:1 EtOAc:hexanes) gave 325 mg of the title compound: MS (ES+) 450.3 (MH+), 472.2 (M+Na).

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b) 3-hydroxy-4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-pyrrolidine hydrochloride

To a solution of the carbamate (310 mg, 0.69 mmol) in dry EtOAc (5.0 mL) was bubbled HCl gas for approximately 5 minutes. The reaction was stirred until TLC analysis indicated the complete consumption of the starting material. The

reaction was then concentrated *in vacuo* to give 249 mg of the title compound: MS (ES+) 350.3 (MH+)

c) 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinol

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To a solution of the amine hydrochloride from the previous step (249 mg, 0.64 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (10 mL) was added CBZ-leucine (170.4 mg, 0.64 mmol), HOBT (86.5 mg, 0.64 mmol), NMM (300 uL) and EDC (147.2 mg, 0.77 mmol). The reaction was allowed to stir at room temperature for 2 hours whereupon it was diluted with ethyl acetate and worked up as described previously. Column chromatography of the residue (3:1EtOAc:hexanes) gave 104 mg of the title compound: MS (ES+) 597.1 (MH+), 619.1 (M+Na).

d) 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

To a 0°C solution of the alcohol (100 mg, 0.17 mmol) in acetone (5.0 mL) was added Jone's reagent dropwise until the brown color persisted. The reaction was allowed to warm to room temperature and stirred approximately 48 hours whereupon it was quenched with isopropanol, diluted with EtOAc and washed sequentially with sat. K<sub>2</sub>CO<sub>3</sub>, water and brine. The organic layer was dried (MgSO<sub>4</sub>), filtered and concentrated. Column chromatography of the residue (3:1 EtOAc:hexanes) gave 31 mg of the title compound: MS (ES+) 595.1 (MH+), 617.0 (M+Na).

## Compound 2. Preparation of 4-IN-I(phenylmethoxy)carbonyl]-L-leucyl]-1-NIN-(methyl)-L-leucyl)]-3-pyrrolidinone

a) 4-{N-{(phenylmethoxy)carbonyl}-L-leucyl}-1-{N-{(tert-butoxy)carbonyl}-N-(methyl)-L-leucyl}-3-pyrrolidinol

To a solution of 3-hydroxy-4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-pyrrolidine (350 mg) was added N-BOC-N-methyl-leucine (222 mg, 0.0.91 mmol), HOBT(122.5 mg, 0.91 mmol), EDC (208.6 mg, 1.08 mmol) and N-methyl morpholine (0.3 mL, 2.72 mmol). The reaction was stirred at room temperature until complete by TLC analysis. Workup and column chromatography (1:1 Hex:EtOAc) gave 480 mg of the title compound which was used in the following reaction: MS (ES+) 477.4, 577.4 (MH+), 599.4 (M+Na).

b) 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(tert-butoxy)carbonyl]-N-(methyl)-L-leucyl]-3-pyrrolidinone

To a -78°C solution of oxalyl chloride (0.11 mL, 1.23 mmol) in CH<sub>2</sub>Cl<sub>2</sub> was added DMSO (0.17 mL, 2.46 mmol) dropwise. The reaction was allowed to stir at -78°C for 20 minutes whereupon a solution of the alcohol (474 mg, 0.82 mmol) in CH<sub>2</sub>Cl<sub>2</sub> was added dropwise. The reaction was stirred at -78°C for 30 minutes whereupon triethylamine (0.57 mL) was added in a single portion and allowed to warm to room temperature. Workup and column chromatography (2:1 hexanes:ethyl acetate) gave 247 mg of the title compound: MS (ES+) 475, 575 (M+H), 597 (M+Na).

- c) 4-[N-[(phenylmethoxy)carbonyi]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone hydrochloride
- To a room temperature solution EtOAc/HCl was added the carbamate. The reaction was stirred until complete by TLC analysis. Concentration gave the title compound: MS (ES+) 475 (M+H, 100%).

Compound 3. Preparation of 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

a) 3-hydroxy-4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-pyrrolidinecarboxylic acid 1,1dimethylethyl ester

- 3-hydroxy-4-amino-1-pyrrolidinecarboxylic acid, 1,1-dimethylethyl ester was coupled with iso-nicotinoyloxycarbonyl leucine in a similar manner as that described above to give 8.5 grams of the title compound: MS (ES+) 451 (MH+, 100%).
- b) 3-hydroxy-4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-pyrrolidine
   30 hydrochloride
   The carbamate from the previous step was deprotected with EtOAc/HCl to give 8.4 grams of the title compound after concentration: MS (ES+)351 (MH+, 100%).
- c) 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-35 [(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinol

To a solution of CBZ leucinal (155 mg) in CH<sub>2</sub>Cl<sub>2</sub> was added triethylamine (0.09 mL) and the amine hydrochloride (200 mg, 0.52 mmol) from the previous step. The reaction was stirred at room temperature for 2 hours whereupon the majority of the solvent was removed *in vacuo*. The mixture was redissolved in CH<sub>2</sub>Cl<sub>2</sub> and sodium triacetoxyborohydride was added. The reaction was stirred at room temperature for 4 hours. Workup and column chromatography (5% methanol/chloroform) gave 200.5 mg of the title compound: MS(ES+) 583 (MH+, 100%).

d) 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone
 To a DMSO (2 mL) solution of the alcohol (50 mg, 0.09 mmol) from the previous
 step was added triethylamine (0.07 mL, 0.52 mmol) and pyridine/sulfur trioxide
 complex (41 mg, 0.26 mmol). The reaction was maintained at room temperature
 until complete by TLC analysis. Workup and chromatography (5%
 methanol/chloroform) gave 37 mg of the title compound: MS (ES+) 582 (MH+,
 100%).

# Compound 4. Preparation of (3S)-3-[(N-benzyloxycarbonyl)-L-leucinyllamino-1-(1-propoxy)-5-methyl-2-hexanone

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(3S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-1-diazo-5-methyl-2-hexanone (150 mg, 0.37 mmol) was dissolved in 1-propanol (2.5 ml), then rhodium acetate (2 mg) was added and the reaction was stirred at RT for 2h. The reaction mixture was chromatographed (silica gel, 20% EtOAc/hexanes) to yield the title compound as a white solid (59 mg, 37%). MS(ES) M+H $^{+}$  = 435, M+ NH $_{4}^{+}$  = 452, 2M+H $^{+}$  = 869.6.

## Compound 5. Preparation of bis-(Cbz-leucinyl)-1.3-diamino-propan-2-one

Cbz-leucine (500 mg, 1.88 mmol), EDCI (558 mg, 1.88 mmol) was dissolved in DMF (4.0 ml) with 1,3-diarnino-propan-2-ol (85 mg, 0.94 mmol) and Hunig's base (0.3 ml, 1.88 mmol) and was stirred at RT overnight. The reaction was diluted with EtOAc (20 ml) and was extracted with water (2 x 20 ml). The combined organics were dried with magnesium sulfate, filtered, concentrated in vacuo. The intermediate was then dissolved in acetone (4.0 ml) and Jones reagent

(2.0 ml, 1.5 M) was added dropwise and the reaction was stirred at RT overnight. The excess Jones reagent was then quenched with isopropanol (1.0 ml), then the reaction was diluted with EtOAc (20 ml) and was extracted with water (2x 20 ml) to remove the inorganic salts. The combined organics were dried with magnesium sulfate, filtered, concentrated, and chromatographed (silica gel, 2-5% MeOH/methylene chloride) to give the title compound as a white solid (410 mg, 75%). MS(ES) M+H\*=583, M+Na\*=605.

# Compound 6. Preparation of 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide

#### a) methyl 3-benzyloxybenzoate

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To a suspension of NaH (0.395 g, 9.87 mmol, 60% in mineral oil) in DMF (20 mL) was added methyl 3-hydroxybenzoate (1.0 g, 6.58 mmol). After stirring for 15 min at room temperature, benzyl bromide (1.1 g, 6.58 mmol) was added. After stirring at room temperature for 3h, the solution was partitioned between ethyl acetate and water. The organic layer was washed with water (2 X 75 mL), saturated aqueous sodium bicarbonate, and brine, then dried (MgSO<sub>4</sub>), filtered and concentrated to yield an off-white solid (1.013 g, 4.2 mmol). <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) d 7.67 (m, 2H), 7.48-7.34 (m. 6H), 7.19 (m, 1H), 5.12 (s, 2H), 3.95 (s, 3H).

#### b) 3-benzyloxybenzoic acid

To a solution of the compound of Example 6(a) (0.400 g, 1.65 mmol) in THF (2 mL) and water (2 mL) was added lithium hydroxide monohydrate (0.076 g, 1.82 mmol). After stirring at reflux for 5 h, the solution was partitioned between ethyl acetate and 3N HCl. The organic layer was washed with brine, dried (MgSO<sub>4</sub>), filtered and concentrated to yield a white solid (0.355 g, 1.56 mmol). <sup>1</sup>H NMR (400 MHz, CD<sub>3</sub>OD) d 7.58 (m, 2H), 7.36-7.24 (m. 6H), 7.10 (m, 1H), 5.04 (s, 2H).

30 c) 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide

Following the procedure of Example A, below, except substituting 3-benzyloxybenzoic acid for N-acetyl-L-leucine and 2-[N-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide for 2-[N-(N-benzyloxycarbonyl-L-alanyl)]carbohydrazide,

the title compound was prepared as a white solid (0.062 g, 25%). MS(ESI): 548.1 (M+H)<sup>+</sup>.

#### Example A

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5 Preparation of 2-[N-(N-acetyl-L-leucinyl)]-2'-[N'-(N-benzyloxycarbonyl-L-alanyl)]carbohydrazide

To a stirring solution of 2-[N-(N-benzyloxycarbonyl-L-alanyl)]carbohydrazide (0.150g, 0.508mmol) in DMF (2mL) was added N-acetyl-L-leucine (0.092g, 0.534mmol), 1-hydroxybenzotriazole (0.014g, 0.102mmol), and 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride (0.102g, 0.534mmol). After stirring at room temperature for 16h, the solution was diluted with ethyl acetate, washed successively with water, saturated aqueous sodium bicarbonate, and brine. The organic layer was dried (MgSO4), filtered and concentrated. The residue was purified by column chromatography (silica gel, methanol/dichloromethane) to yield the title compound as a white solid (0.028 g, 12%). MS(ESI): 451.1 (M+H)+.

# Compound 7. Preparation of (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide

20 a) N-tert-butoxycarbonyl-(L)-leucinamide

To a solution of N-tert-butoxycarbonyl-(L)-leucine (7.0g, 28.1mmol) in dry THF (100mL) at -40°C was added isobutylchloroformate (3.8g, 28.1mmol) and N-methylmorphiline (6.0, 59mmol). After 15 minutes of stirring, ammonia was bubbled through the mixture for an additional 15 minutes, then warmed to room temperature and allowed to stir for 2 hours. Mixture filtered and filtrate concentrated in vacuo to yield title compound as a white solid (6.5, 28.0mmol). 'HNMR (400MHz, CDCl<sub>3</sub>) d 6.38 (br s, 1H), 5.79 (br s, 1H), 5.04 (br d, 1H), 4.13 (m, 1H), 1.71-1.49 (m, 3H), 1.39 (s, 9H), 0.92 (dd, 6H).

30 b) N-tert-butoxycarbonyl-(L)-leucinethioamide

To a stirring solution of the compound of Example 7(a) (6.5, 28.0 mmol) in dry THF was added Lawesson's reagent (6.8g, 16.9 mmol) and the mixture was stirred at room temperature under argon overnight. The solvent was evaporated and the residue chromatographed (silica gel, 12% ethyl acetate/hexane) to give the title compound as a white solid (5.4g, 77%). HNMR (400MHz, CDCl<sub>3</sub>) d 8.54 (br s,

1H), 7.97 (br s, 1H), 5.28 (br d, 1H), 4.52 (m, 1H), 1.72-1.58 (m, 3H), 1.40 (s, 9H), 0.92 (m, 6H).

c) (1S)-1-(tert-butoxycarbonyl)amino-1-(4-carboethoxythiazol-2-yl)-3-methylbutane

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The compound of Example 7(b) (5.4g, 21.7 mmol) was stirred in dry acetone (100mL) under argon at -10°C. Ethylbromopyruvate (4.7g, 23.9mmol) was added and stirred for 1h at -10°C. The solution was poured into a well stirred mixture of chloroform and water and then into saturated sodium bicarbonate solution. The organic phase was separated and the aqueous layer extracted with chloroform. The combined organic extracts were dried over MgSO<sub>4</sub>, filtered and concentrated to an oil. The oily residue was treated with TFAA (5.0g, 23.9mmol) and pyridine (3.8g, 47.8mmol) in dichloromethane for 1h at -20°C. Excess solvent was removed in vacuo and the residue was dissolved in dichloromethane. The solution was washed with saturated aqueous sodium bicarbonate and 1.0N KHSO<sub>4</sub> until pH 7. The solution was dried over magnesium sulfate, filtered and concentrated to an oil which was chromatographed (silica gel, 7.5% ethyl acetate/hexane) to give the title compound as a tan solid (4.5g, 61%). 'HNMR (400MHz, CDCl<sub>3</sub>) d 7.98 (s, 1H), 5.04 (br d, 1H), 4.95 (m, 1H), 4.31 (q, 2H), 1.88 (m, 1H), 1.63 (m, 2H), 1.40 (s, 9H),1.32 (t, 3H), 0.85 (dd, 6H).

d) (1S)-1-(Benzyloxycarbonyl)amino-1-(4-carboethoxythiazol-2-yl)-3-methylbutane

The compound of Example 7(c) (0.250g, 0.731mmol) was dissolved in TFA

(2mL) and stirred at room temperature for 15 minutes when diluted with methanol and concentrated in vacuo. The residue was dissolved in methylene chloride and treated with triethylamine (0.739g, 7.31mmol) followed by benzyl chloroformate (1.2g, 7.31mmol). The solution stirred at room temperature for 2h when partition between ethyl acetate/water. The organic layer was washed with brine, collected, dried (MgSO<sub>4</sub>) and concentrated to a residue that was chromatographed (silica gel, 15% ethyl acetate/hexane) to give the title compound as an oil (0.198g, 72%).

HNMR (400MHz, CDCl<sub>3</sub>) d 8.01 (s, 1H), 7.32 (m, 5H), 5.51 (br d, 1H), 5.14 (m, 1H), 5.10 (s, 2H), 4.37 (q, 2H), 1.93 (m, 1H), 1.81-1.67 (m, 2H), 1.39 (t, 3H), 0.95 (m, 6H).

e) (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'- (N-benzyloxycarbonyl-L-leucinyl)hydrazide

Following the procedure of Example B(a)-(d), below, except substituting (1S)-1-(Benzyloxycarbonyl)amino-1-(4-carboethoxythiazol-2-yl)-3-methylbutane for (1S)-1-benzyloxycarbonylamino-1-(2-carboethoxythiazol-4-yl)-3-methylbutane in step (c), the title compound was prepared. MS (MH\*): 610.0

#### Example B

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Preparation of (1S.2'R)-N-4-[[(1-benzyloxycarbonyl)aminol-3-methylbutyl]thiazol-2-ylcarbonyl-N'-2'-(benzyloxycarbonyl)amino-4'-methylpentanoylhydrazide

a) N-benzyloxycarbonyl-L-leucinyl bromomethyl ketone

1-methyl-3-nitro-1-nitrosoguanidine (6.65 g, 45.2 mmol) in ether (225 mL) is cooled to 0°C. 40% sodium hydroxide is added slowly and the diazomethane is allowed to collect in the ether solution for 30 minutes at 0°C. The ether solution is then decanted and left at 0 °C.

N-Cbz-L-leucine (2.10 g, 7.6 mmol) was dissolved in THF (10 mL), cooled to -40 °C, and 4-methylmorpholine (0.77 g, 7.6 mmol, 0.83 mL) was added, followed by dropwise addition of isobutyl chloroformate (1.04 g, 7.6 mmol, 0.98 mL). After 15 min, the solution was filtered into the previously prepared 0 °C solution of ethereal diazomethane. The resulting solution was allowed to stand at 0 °C for 23 h. HBr (30% in acetic acid) (45.2 mmol, 9 mL) was added and the resulting solution was stirred at 0 °C for 5 min, then washed sequentially with 0.1 N HCl, saturated aqueous NaHCO3 and saturated brine, then dried (MgSO4), filtered and concentrated to give the title compound as a colorless oil (2.43 g, 94%).

b) (1S)-1-benzyloxycarbonylamino-1-(2-carboethoxythiazol-4-yl)-3-methylbutane
A solution of the compound of Example B(a) (1.57 g, 4.58 mmol) and ethyl
thiooxamate (0.61 g, 4.58 mmol) in ethanol (10 mL) was heated at reflux for 4 h.
The solution was then cooled, concentrated and the residue was purified by flash
chromatography on 230-400 mesh silica gel. eluting with 1:4 ethyl acetate/hexanes,
to give the title compound as a yellow oil (1.0 g, 58%). 1H NMR (400 MHz,
CDC13) d 7.41 (s, 1H), 7.34-7.31 (m, 5H), 5.40 (d, 1H), 5.10 (d, 1H), 5.05 (d, 1H),

4.98 (q, 1H), 4.48 (q, 2H), 1.80-1.76 (m, 2H), 1.57-1.53 (m, 1H), 1.44 (t, 3H), 0.95 (d, 3H), 0.93 (d, 3H).

c) (1S)-1-benzyloxycarbonylamino-1-(2-hydrazinocarbonylthiazol-4-yl)-3-methylbutane

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A solution of the compound of Example B(b) (0.30 g, 0.8 mmol) and hydrazine hydrate (0.40 g, 8.0 mmol, 0.39 mL) in ethanol (8 mL) was allowed to stir at room temperature for 2 h. The solution was then concentrated to yield the title compound as a white foam (0.28 g, 98%). 1H NMR (400 MHz, CDCl3) d 8.29 (s, 1H), 7.37-7.35 (m, 5H), 5.18 (d, 1H), 5.09 (dd, 2H), 4.95 (q, 1H), 4.07 (d, 2H), 1.71 (t, 2H), 1.55 (m, 1H), 0.96 (d, 3H), 0.94 (d, 3H).

d) (1S,2'R)-N-4-[[(1-benzyloxycarbonyl)amino]-3-methylbutyl]thiazol-2-ylcarbonyl-N'-2'-(benzyloxycarbonyl)amino-4'-methylpentanoylhydrazide

A solution of the compound of Example B(c) (100 mg, 0.28 mmol), N-Cbz-L-leucine (80.5 mg, 0.30 mmol), 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride (58.2 mg, 0.30 mmol) and 1-hydroxybenzotriazole (7.5 mg, 0.06 mmol) in DMF (0.6 mmol) was allowed to stir at room temperature for 18 h. The solution was diluted with ethyl acetate and washed successively with water, 0.1 N HCl, saturated aqueous NaHCO3 and saturated brine, then dried (MgSO4), filtered and concentrated. The residue was purified by flash chromatography on 230-400 mesh silica gel, eluting with 1:1 ethyl acetate/hexanes, to provide the title compound as a white solid (111.4 mg, 66%). mp 110-112 °C.

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## Compound 8. Preparation of 2.2'-N.N'-bis-benzyloxycarbonyl-Lleucinylcarbohydrazide

To a stirring solution of N-Cbz-L-leucine (Chemical Dynamics Corp.) (2.94 g, 11.1 mmol) in 22 mL of DMF was added carbohydrazide (0.5 g, 5.6 mmol), 1-(3-5 dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride (2.13 g, 11.1 mmol) and 1-hydroxybenzotriazole (0.3 g, 2.2 mmol). After stirring at room temperature for 22 h, the solution was poured into 500 mL of water. The precipitate was collected by vacuum filtration and washed with water (4 X 150 mL) and dichloromethane (4 X 150 mL), then dried under vacuum to provide the title compound as a white solid 10 (1.49 g, 46%). MS(ESI): 607.1 (M+Na)\*.

## Compound 9. Preparation of 1-N-( N-imidazole acetyl-leucinyl)-amino-3-N-(4phenoxy-phenyl-sulfonyl)-amino-propan-2-one

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1-N-( N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy phenyl sulfonyl)a) amino-propan-2-one

Following the procedure of Example C(a)-(d), below, substituting "imidazole acetic acid" for "4-pyridyl acetic acid", the title compound was prepared: MS(ES) M  $+H^{+}=542.$ 

#### Example C

Preparation of 1-N-(N-Cbz-leucinyl)-amino-3-N-(2-pyridyl-sulfonyl)-aminopropan-2-one

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- 1-N-(N-Cbz-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-aminoa) propan-2-ol
- 1,3-Diamino propan-2-ol (6.75 g, 75 mmol) was dissolved in DMF (100ml) and Cbz-leucine (20g, 75.5 mmol), HOBT-hydrate (11g, 81.5 mmol), and EDCI (15.5g, 81.2 mmol) were added. The reaction was stirred overnight at RT. A 30 portion of the reaction mixture (30 ml) was concentrated in vacuo, then ether (50 ml) and MeOH (30 ml) were added. A 1N solution of hydrochloric acid in ether was added (1 M, 30 ml) and a white gum formed, which was washed several times with ether. MeOH-acetone were added and heated until the gum became a white solid. The white solid was dissolved in DMF (25 ml) and DIEA (5ml), then 4-phenoxy

phenyl sulfonyl chloride was added. The reaction was stirred for 2h, concentrated in vacuo, then chromatographed (silica gel, 1:1 EtOAc: hexanes) to provide the desired product as a white solid.

- b) Leucinyl-amino-3-N-(4-phenoxy phenyl sulfonyl)-amino-propan-2-ol
  1-N-(Cbz-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan2-ol (1.0g, 1.8 mmol) was dissolved in EtOH (30 ml), then 10% Pd/C (0.22g) was added followed by 6N hydrochloric acid (2.5 ml), and the reaction was stirred under a balloon of hydrogen gas for 4h at RT. The reaction mixture was filtered,
   concentrated, and azeotroped with toluene to provide a white glass which was used in the next reaction without further purification.
  - c) 1-N-(N-4-pyridyl acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-ol

Leucinyl-amino-3-N-(4-phenoxy phenyl sulfonyl)-amino-propan-2-ol (0.36 g, 0.76 mmol) was dissolved in DMF (5 ml), then NMM (0.45 ml, 4 mmol) was added followed by 4-pyridyl acetic acid (0.13g, 0.75 mmol) and HBTU (0.29g, 0.76 mmol) and the reaction was stirred at RT overnight. The reaction mixture was concentrated in vacuo, then chromatographed (silica gel, 5%MeOH: methylene chloride) to provide the desired product as a white solid (90 mg, MS(ES): M+H+ = 555.

d) 1-N-(N-4-pyridyl acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one

1-N-(N-4-pyridyl-acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-ol (45 mg, 0.08 mmol) was dissolved in acetone (5ml), then 1N hydrochloric acid (2 ml) was added. The reaction was concentrated in vacuo, then redissolved in acetone. Jones reagent (1.5 M, several drops) was added and the reaction mixture was stirred for 6h at RT. Isopropanol (0.5 ml) was added and the reaction mixture was concentrated in vacuo. The reaction was diluted with pH 7 buffer and then was extracted with EtOAc, dried with magnesium sulfate, filtered, concentrated in vacuo, then chromatographed (silica gel, 5% MeOH-methylene chloride) to give the desired product as a white solid (27 mg, 50%): MS(ES): M+H+ = 553.

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#### Crystallization of the protein and protein-inhibitor complexes B.

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Human cathepsin K was expressed in baculovirus cells for the first eight of the nine inhibitors described below. Conditioned media containing expressed pro-cathepsin K was loaded directly onto an S-Sepharose column pre-equilibrated with 25 mM phosphate buffer at pH 8. The column was eluted with a NaCl gradient. Fractions containing pro-cathepsin K were pooled, concentrated to 2.5 mg/ml and activated to mature cathepsin K in 50 mM sodium acetate buffer pH 4.0 containing 20 mM L-cysteine and 1% mature cathepsin K as seed. The activation was monitored using CBZ-Phe-Arg-AMC, as fluorogenic substrate and by SDS-PAGE. When the increasing specific activity reached a plateau (ca. 15 µmol/min/mg), the reaction was stopped by the addition of inhibitor. The inhibited mature cathepsin K was concentrated and dialyzed against 20 mM MES, 50 mM NaCl, 2 mM Lcysteine, pH 6.

Protein preparation for cathepsin K complex with 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone (only)

Human cathepsin K was expressed in E. coli. The cell pellet from 1 L of bacterial culture weighing 2.35 gm. was washed with 50 mL of 50 mM Tris/HCl, 5 mM EDTA, 150 mM NaCl, pH 8.0. After centrifugation at 13,000 x g for 15 mins, the washed pellet was resuspended into 25 mL of the same buffer prepared at 4° C and lysed by passage twice through a cell disruptor (Avestin) at 10,000 psi. The lysate was centrifuged as above, the supernatant decanted and the pellet suspended in 25 mL 50 mM Tris/HCl, 10 mM DTT, 5 mM EDTA, 150 mM NaCl, pH 8.0 25 containing either 8 M urea or 6 M guanidine HCl. After stirring at 4° C for 30 mins, insoluble cellular debris was removed by centrifugation at 23,000 x g for 30 mins and the supernatant clarified by filtration (0.45 um, Millipore).

Varying amounts of the proenzyme form of cathepsin K were refolded by quick dilution into stirring, N2 (g) sparged 50 mM Tris/HCl, 5 mM EDTA, 10 mM reduced and 1 mM oxidized glutathione, 0.7 M L-arginine pH 8.0 and stirred overnight at 4° C. After concentration to ca.1 mg/mL using a stirred cell fitted with a YM-10 membrane (Amicon), the sample was clarified by centrifugation and filtration then dialyzed against 25 mM Na<sub>2</sub>PO<sub>4</sub>, 1.0 M NaCl, pH 7.0. The dialysate was applied at a LFR= 23 cm/hr to

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a 2.6 x 90 cm column of Superdex 75 (Pharmacia) pre-equilibrated in 25 mM Na<sub>2</sub>PO<sub>4</sub>, 1.0 M NaCl, pH 7.0. The cathepsin K proenzyme was pooled based upon purity as observed on a reduced, SDS-PAGE gel.

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Crystals of mature activated cathepsin K complexed with inhibitor grew to a size of approximately 0.2 mm3 in about six days at 20°C. The concentration of inhibited cathepsin K used in the crystallization was approximately 8 mg./ml. The method of vapor diffusion in hanging drops was used to grow crystals from the solution of cathepsin K - inhibitor complex. The initial crystal structure to be determined was that of cathepsin K in complex with the cysteine protease inhibitor E64. Crystals of mature activated cathepsin K complexed with E-64 grew to a size of approximately 0.2 mm<sup>3</sup> in six days at 20°C. The concentration of E-64-inhibited cathepsin K used in the crystallization was 8 mg/ml. Vapor diffusion was used in hanging drops from a solution of 10% PEG 8000, 0.1 M Na+/K+ phosphate at pH 6.2 containing 0.2M NaCl. Crystals of the complex are orthorhombic, space group P2,2,2, with cell constants of a=38.4, b=50.7, and c=104.9 Angstroms. This crystal form will be referred to as Form II. The crystals contain one molecule in the asymmetric unit and contain approximately 40% solvent with a Vm value of 2.1 A<sup>3</sup>/Dalton. X-ray diffraction data were measured from a single crystal using a Siemens two-dimensional position-sensitive detector on a Siemens rotating anode generate operating a 5 KW. The structure was determined by molecular replacement using X-PLOR. The starting model consisted of all atoms of the main chain of papain and those side chain atoms predicted to be homologous between the two proteins as determined from sequence alignment. The cross rotation function was calculated using x-ray diffraction data from 10 to 4 Å and a radius of integration of 32 Å. The highest peak was 6.0  $\sigma$ . A translation search was carried out using data from 8 to 3.5 Ångstroms resulting in the highest peak of 12.5  $\sigma$ . The resulting model gave an R<sub>c</sub> factor of 0.488. This model was refined by rigid-body refinement, and the resulting phases were used to calculate Fourier maps with coefficients IFo-FcI and 12F<sub>0</sub>-F<sub>c</sub>l, into which the atomic model of cathepsin K was built using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building. The structure was refined using X-PLOR. The electron density for E-64 was clear in the maps. The inhibitor was built into density and several additional cycles of map fitting and refinement were carried out to a final Rc of 0.191.

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Crystallization of the complex of cathepsin K with 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyllamino-5-methyl-1-(1-propoxy)-2-hexanone

Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 10% isopropanol, 0.1 M NaPO<sub>4</sub> / citrate at pH 4.2. Crystals of the complex are tetragonal, space group P43212, with cell constants of a=57.6 Å, and c=131.2 Å. This crystal form will be referred to as Form III. Diffraction data were collected as described above. The crystals contain one molecule in the asymmetric unit and contain 36% solvent with a  $V_m$  value of 2.3 Å  $^3$ /Dalton. The structure was determined by molecular replacement using X-PLOR at 2.5 Ångstroms resolution. The starting model consisted of all protein atoms of the orthorhombic form of cathepsin K-E64 structure. Molecular replacement was carried out as described above for the cathepsin K-E64 structure determination. The model was refined by rigid-body refinement using X-PLOR, and the resulting phases were used to calculate Fourier maps with coefficients |Fo-Fc| and |2Fo-Fc|, into which the atomic model of the inhibitor was built using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building. The structure was refined using X-PLOR. Several cycles of map fitting and refinement were carried out to a final  $R_{\rm C}$  of 0.245.

Crystallization of the complex of cathepsin K with 2-[N-(3-benzyloxybenzoyl)]-2'[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide

Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 22.5% PEG 8000, 0.075 M sodium acetate at pH 4.5 containing 0.15 M Li<sub>2</sub>SO<sub>4</sub>. Crystals of the complex grew as Form III. Diffraction data were collected as described above. The structure was determined by rigid body refinement with X-PLOR utilizing the previous Form III protein model at 2.4 Ångstroms resolution.
 Fourier maps with coefficients IF<sub>0</sub>-F<sub>c</sub>I and I2F<sub>0</sub>-F<sub>c</sub>I were used to fit the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement (X-PLOR) was used to refine the structure during model building. Several cycles of map fitting and refinement were carried out to a final R<sub>c</sub> of 0.237.

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Crystallization of the complex of cathepsin K with bis-(Cbz-leucinyl)-1.3-diamino-propan-2-one

Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 10% isopropanol, 0.1 M NaPO<sub>4</sub> / citrate at pH 4.2. Crystals of the complex grow as Form III. Diffraction data were collected as described above. The structure was determined by rigid body refinement of the previous Form III protein model at 2.6 Ångstroms resolution. Fourier maps with coefficients IF<sub>0</sub>-F<sub>c</sub>I and I2F<sub>0</sub>-F<sub>c</sub>I were used to fit the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building. Several cycles of map fitting and refinement were carried out using X-PLOR to a final R<sub>c</sub> of 0.210.

Crystallization of the complex of cathepsin K with 4-IN
[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone

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Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution 18% PEG 8000, 0.6 M sodium acetate at pH 4.5 containing 0.12 M Li<sub>2</sub>SO<sub>4</sub>. Crystals of the complex grow in Form III. Diffraction data were collected as described above. The structure was determined by rigid body refinement of the previous Form III protein model with X-PLOR at 2.4 Ångstroms resolution. Fourier maps with coefficients  $|F_0-F_c|$  and  $|2F_0-F_c|$ , were used to the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building using X-PLOR. Several cycles of map fitting and refinement were carried out to a final  $R_c$  of 0.218.

Crystallization of the complex of cathepsin K with (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl)thiazol-4-ylcarbonyll-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide

Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 30% MPD, 0.1 M MES at pH 7.0 and 0.1 M tris buffer at pH 7.0. Crystals of the complex are Form II. Diffraction data were collected as described above. The structure was determined by rigid body refinement of the previous Form II protein model with X-PLOR at 2.3 Ångstroms resolution. Fourier maps with

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coefficients  $|F_0-F_C|$  and  $|2F_0-F_C|$ , were used to the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building using X-PLOR. Several cycles of map fitting and refinement were carried out to a final  $R_C$  of 0.211.

Crystallization of the complex of cathepsin K with 2.2'-N.N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide

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Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 33% MPD, 0.1 M MES at pH 7. Crystals of the complex grow as Form II. Diffraction data were collected as described above. The structure was determined by rigid body refinement of the previous Form II protein model with X-PLOR at 2.2 Ångstroms resolution.. Fourier maps with coefficients IF<sub>0</sub>-F<sub>c</sub>I and I2F<sub>0</sub>-F<sub>c</sub>I, were used to the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building using X-PLOR. Several cycles of map fitting and refinement were carried out to a final R<sub>c</sub> of 0.208.

Crystallization of the complex of cathepsin K with 4-IN
[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3
pyrrolidinone

Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 28% MPD, 0.1 M MES at pH 7.0 and 0.1 M tris buffer at pH 7.0.

Crystals of the complex Form II. Diffraction data were collected as described above. The structure was determined by rigid body refinement of the previous Form II protein model with X-PLOR at 2.3 Ångstroms resolution. Fourier maps with coefficients IF<sub>0</sub>-F<sub>c</sub>I and I2F<sub>0</sub>-F<sub>c</sub>I, were used to the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building using X-PLOR. Several cycles of map fitting and refinement were carried out to a final R<sub>c</sub> of 0.193.

Crystallization of the complex of cathensin K with 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

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Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 30% MPD, 0.1 M MES at pH 7.0 and 0.1 M tris buffer at pH 7.0. Crystals of the complex Form II. Diffraction data were collected as described above.

The structure was determined by rigid body refinement of the previous Form II protein model with X-PLOR at 2.2 Ångstroms resolution.. Fourier maps with coefficients  $|F_0-F_c|$  and  $|2F_0-F_c|$ , were used to the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building using X-PLOR. Several cycles of map fitting and refinement were carried out to a final R<sub>C</sub> of 0.267. 10

Crystallization of the complex of cathepsin K with 1-N-(N-imidazole acetylleucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one

Crystals of mature activated cathepsin K complexed with the inhibitor grew from a solution of 18% PEG 8000, 0.6 M sodium acetate at pH 4.5 containing 0.12 M Li<sub>2</sub>SO<sub>4</sub>. Crystals of the complex are Form III. Diffraction data were collected as described above. The structure was determined by rigid body refinement of the previous Form II protein model at 2.5 Ångstroms resolution.. Fourier maps with coefficients IF<sub>0</sub>-F<sub>c</sub>I and I2F<sub>0</sub>-F<sub>c</sub>I were used to fit the atomic model of the inhibitor using the molecular graphics program FRODO. Conventional positional refinement was used to refine the structure during model building. Several cycles of map fitting and refinement were carried out using X-PLOR to a final R<sub>C</sub> of 0.246. **Abbreviations** 

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E-64, [1-[N-[(L-3-trans-carboxyoxirane-2carbonyl)-L-leucyl]amino]-4-guanidinobutane] CBZ, benzyloxycarbonyl AMC, aminomethylcoumarin MPD, 2 methyl-2,4-pentanediol PIPES, piperazone-N,N-bis(2-ethanesulfonic acid) MES, 2-(N-morpholino)-ethanesulfonic acid

tris, tris(hydroxymethyl)-aminomethane

PEG, polyethyleneglycol

M. Molar

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 $R_c = \Sigma I(F_o - F_c)I/F_o$ 

 $F_0$  = observed structure amplitude

 $F_c$  = calculated structure amplitude

EDTA, ethylenediaminetetraacetic acid

DTT, 1,4-dithiothreitol

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SDS-PAGE, sodium dodecylsulfate polyacrylamide gel electrophoresis

This invention is not to be limited in scope by the specific embodiments described herein. Indeed, various modifications of the invention in addition to those described herein will become apparent to those skilled in the art from the foregoing description. Such modifications are intended to fall within the scope of the appended claims.

The disclosures of the patents, patent applications and publications cited herein are incorporated by reference in their entireties.

TABLE I

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (Å<sup>2</sup>) for cathepsin K.

Residue Atom	x	Y	Z	В
1 ALA N	-3.94	11.01	90.45	15.00
1 ALA CA	-4.70	12.30	90.45	15.00
1 ALA C	-4.40	13.14	89.20	15.00
1 ALA O	-3.34	12.99	88.57	15.00
1 ALA CB	-4.36	13.12	91.73	15.00
2 PRO N	-5.36	14.01	88.80	15.00
2 PRO CA	-5.19	14.86	87.61	15.00
2 PRO C	-4.35	16.05	88.06	15.00
2 PRO O	-4.89	17.00	88.65	15.00
2 PRO CB	-6.62	15.33	87.31	15.00
2 PRO CG	-7.53	14.58	88.32	15.00
2 PRO CD	-6.63	14.31	89.47	15.00
3 ASP N	-3.04	16.00	87.87	15.00
3 ASP CA	-2.25	17.14	88.30	15.00
3 ASP C	-2.27	18.18	87.20	15.00
3 ASP O	-1.57	18.02	86.20	15.00
3 ASP CB	-0.82	16.75	88.67	15.00
3 ASP CG	-0.09	17.85	89.45	15.00
3 ASP OD1	-0.74	18.83	89.89	15.00
3 ASP OD2	1.14	17.73	89.63	15.00
4 SER N	-3.10	19.21	87.36	15.00 15.00
4 SER CA	-3.19	20.26	86.35 86.77	15.00
4 SER C	-3.97	21.51	87.48	15.00
4 SER O	-4.97	21.44 19.72	85.03	15.00
4 SER CB	-3.77 5.17	19.72	85.08	15.00
4 SER OG	-5.17 -3.50	22.65	86.28	15.00
5 VAL N 5 VAL CA	-4.10	23.94	86.54	15.00
5 VAL CA 5 VAL C	-4.27	24.65	85.17	15.00
5 VAL C	-3.43	24.48	84.28	15.00
5 VAL CB	-3.22	24.79	87.51	15.00
5 VAL CG1	-1.80	24.88	87.00	15.00
5 VAL CG2	-3.79	26.17	87.69	15.00
6 ASP N	-5.39	25.34	84.99	15.00
6 ASP CA	-5.67	26.08	83.76	15.00
6 ASP C	-6.40	27.34	84.22	15.00
6 ASP O	-7.63	27.33	84.43	15.00
6 ASP CB	-6.55	25.25	82.82	15.00
6 ASP CG	-6.81	25.95	81.48	15.00

		Indus.		
6 ASP OD1	-6.11	26.94	81.14	15.00
6 ASP OD2	-7.72	25.49	80.75	15.00
7 TYR N	-5.64	28.42	84.37	15.00
7 TYR CA	-6.15	29.70	84.84	15.00
7 TYR C	-7.18	30.35	83.96	15.00
7 TYR O	-7.76	31.36	84.33	15.00
7 TYR CB	-5.00	30.67	85.09	15.00
7 TYR CG	-4.06	30.20	86.18	15.00
7 TYR CD1	-4.41	30.29	87.52	15.00
7 TYR CD2	-2.82	29.64	85.86	15.00
7 TYR CE1	-3.55	29.86	88.52	15.00
7 TYR CE2	-1.96	29.21	86.84	15.00
7 TYR CZ	-2.33	29.31	88.17	15.00
7 TYR OH	-1.48	28.86	89.14	15.00
8 ARG N	-7.41	29.78	82.79	15.00
8 ARG CA	-8.41	30.30	81.87	15.00
8 ARG C	-9.77	30.07	82.53	15.00
8 ARG O	-10.65	30.93	82.49	15.00
8 ARG CB	-8.33	29.58	80.53	15.00
8 ARG CG	-7.00	29.76	79.85	15.00
8 ARG CD	-7.00	29.12	78.48	15.00
8 ARG NE	-7.27	27.69	78.52	15.00
8 ARG CZ	-6.85	26.83	77.58	15.00
8 ARG NH1	-6.15	27.27	76.54	15.00
8 ARG NH2	-7.15	25.54	77.69	15.00
9 LYS N	-9.90	28.94	83.20	15.00
9 LYS CA	-11.12	28.60	83.91	15.00
9 LYS C	-11.16	29.33	85.28	15.00
9 LYS O	-11.96	28.99	86.15	15.00
9 LYS CB	-11.18	27.08	84.13	15.00
9 LYS CG	-11.04	26.25	82.86	15.00 15.00
9 LYS CD	-11.09	24.72	83.11	15.00
9 LYS CE	-9.80	24.15	83.76	15.00
9 LYS NZ	-9.78	22.65	83.99	15.00
10 LYS N	-10.33	30.35	85. <b>4</b> 7	15.00
10 LYS CA	-10.28	31.03	86.76	15.00
10 LYS C	-10.23	32.55	86.69	15.00
10 LYS O	-10.11	33.20	87.73	15.00
10 LYS CB	-9.10	30.51	87.58	15.00
10 LYS CG	-9.05	28.98	87.72 88.13	15.00
10 LYS CD	-7.68	28.45		15.00
10 LYS CE	-7.54	28.31	89.63	15.00
10 LYS NZ	-7.61	29.62	90.36	15.00
11 GLY N	-10.29	33.11	85.48	٠٠٠٠

11	GLY	CA	-10.27	34.56	85.31	15.00
11	GLY	С	-8.96	35.28	85.53	15.00
11	GLY	0	-8.93	36.49	85.77	15.00
12	TYR	N	-7.86	34.54	85.44	15.00
12	TYR	CA	-6.54	35.11	85.64	15.00
12	TYR	С	-5.97	35.67	84.36	15.00
12	TYR	0	-5.13	36.58	84.39	15.00
12	TYR	CB	-5.57	34.04	86.13	15.00
12	TYR	CG	-5.76	33.63	87.56	15.00
12	TYR	CD1	-6.85	32.86	87.95	15.00
12	TYR	CD2	-4.82	33.98	88.52	15.00
12	TYR	CE1	-7.00	32.46	89.25	15.00
12	TYR	CE2	-4.96	33.58	89.83	15.00
12	TYR	CZ	-6.04	32.81	90.19	15.00
12	TYR	OH	-6.16	32.38	91.49	15.00
13	VAL	N	-6.40	35.09	83.24	15.00
13	VAL	CA	-5.92	35.47	81.92	15.00
13	VAL	C	-6.95	36.24	81.09	15.00
13	VAL	0	-8.15	36.03	81.21	15.00
13	VAL	CB	-5.41	34.21	81.15	15.00
13	VAL	CG1	-6.54	33.26	80.89	15.00
13	VAL	CG2	-4.73	34.61	79.86	15.00
14	THR	N	-6.45	37.19	80.31	15.00
14	THR	CA	-7.27	38.02	79.44	15.00
14	THR	С	-7.39	37.38	78.05	15.00
14	THR	0	-6.69	36.41	77.74	15.00
14	THR	CB	-6.63	39.42	79.32	15.00
14	THR	OG1	-5.21	39.28	79.27	15.00
14	THR	CG2	-7.00	40.28	80.52	15.00
15	PRO	N	-8.31	37.88	77.20	15.00
15	PRO	CA	-8.50	37.34	75.86	15.00
	PRO		-7.23	37.45	75.01	15.00
15	PRO		-6.38	38.30	75.30	15.00
	PRO		-9.61	38.22	75.30	15.00
	PRO		-10.38	38.60	76.51	15.00
	PRO		-9.28	38.96	77.45	15.00
	VAL		-7.14	36.65	73.95	15.00
	VAL		-5.97	36.64	73.08	15.00
	VAL		-5.86	37.87	72.18	15.00
	VAL		-6.80	38.23	71.47	15.00
16			-5.94	35.38	72.22	15.00
	VAL		-4.70	35.38	71.34	15.00
16	VAL	CG2	-5.97	34.18	73.10	15.00
17	LYS	N	-4.70	38.52	72.22	15.00

17	LYS	CA	-4.47	39.71	71.43	15.00
17	LYS	С	-3.49	39.39	70.31	15.00
17	LYS	0	-2.82	38.36	70.34	15.00
17	LYS	CB	-3.93	40.85	72.30	15.00
17	LYS	CG	-4.99	41.75	72.95	15.00
17	LYS	CD	-5.84	41.01	73.99	15.00
17	LYS	CE	-5.88	41.72	75.34	15.00
17	LYS	NZ	-4.53	41.86	75.97	15.00
18	ASN	N	-3.43	40.30	69.35	15.00
18	ASN	CA	-2.58	40.20	68.17	15.00
18	ASN	С	-1.54	41.29	68.24	15.00
18	ASN	0	-1.89	42.46	68.35	15.00
18	ASN	CB	-3.42	40.41	66.91	15.00
18	ASN	CG	-2.71	39.96	65.64	15.00
18	ASN	OD1	-1.74	40.58	65.19	15.00
18	ASN	ND2	-3.19	38.87	65.06	15.00
19	GLN	N	-0.27	40.92	68.15	15.00
19	GLN	CA	0.79	41.91	68.23	15.00
19	GLN	С	0.97	42.67	66.95	15.00
19	GLN	0	1.54	43.76	66.94	15.00
19	GLN	СВ	2.10	41.26	68.59	15.00
19	GLN	CG	2.54	40.22	67.63	15.00
19	GLN	CD	3.88	39.71	67.99	15.00
19	GLN	OE1	4.04	38.93	68.92	15.00
19	GLN	NE2	4.89	40.20	67.31	15.00
20	GLY	N	0.51	42.07	65.86	15.00
20	GLY	CA	0.62	42.69	64.56	15.00
20	GLY	С	1.98	42.49	63.90	15.00
20	GLY	0	2.53	41.39	63.86	15.00
21	GLN	N	2.50	43.60	63.38	15.00
21	GLN	CA	3.77	43.62	62.67	15.00
21	GLN	С	4.94	43.82	63.62	15.00
21	GLN	0	6.05	43.35	63.36	15.00
21	GLN	CB	3.73	44.74	61.64	15.00
21	GLN	CG	2.68	44.55	60.56	15.00
21	GLN	CD	2.88	43.25	59.82	15.00
21	GLN	OE1	4.01	42.87	59.52	15.00
21	GLN	NE2	1.79	42.55	59.56	15.00
22	CYS	N	4.68	44.56	64.69	15.00
22	CYS	CA	5.65	44.87	65.73	15.00
22	CYS	С	6.15	43.60	66.46	15.00
	CYS		5.37	42.68	66.72	15.00
22	CYS	CB	4.97	45.82	66.71	15.00
22	CYS	SG	5.96	46.40	68.11	15.00

23	GLY	N	7.45	43.53	66.73	15.00
23	GLY	CA	8.00	42.37	67.42	15.00
23	GLY	C	7.84	42.62	68.90	15.00
23	GLY	0	8.80	42.94	69.61	15.00
24	SER	N	6.60	42.54	69.37	15.00
24	SER	CA	6.33	42.80	70.77	15.00
24	SER	С	5.75	41.63	71.54	15.00
24	SER	0	4.72	41.77	72.20	15.00
24	SER	СВ	5.44	44.05	70.90	15.00
24	SER	OG	4.14	43.81	70.40	15.00
25	CYC	N	6.39	40.47	71.48	15.00
25	CYC	CA	5.88	39.34	72.25	15.00
25	CYC	CB	6.32	38.03	71.63	15.00
25	CYC	SG	8.04	38.06	71.17	15.00
25	CYC	C	6.33	39.44	73.72	15.00
25	CYC	0	5.67	38.91	74.62	15.00
25	CYC	01	7.96	38.09	69.44	15.00
26	TRP		7.45	40.14	73.95	15.00
26	TRP	CA	7.97	40.33	75.30	15.00
26	TRP	C	7.04	41.24	76.05	15.00
26	TRP	0	6.67	40.96	77.18	15.00
26	TRP		9.37	40.92	75.27	15.00
26	TRP	CG	9.47	42.24	74.61	15.00
26	TRP	CD1	9.74	42.49	73.30	15.00
26	TRP	CD2	9.32	43.52	75.24	15.00
26	TRP		9.78	43.84	73.06	15.00
26	TRP		9.52	44.50	74.24	15.00
26	TRP		9.04	43.94	76.55	15.00
26	TRP		9.45	45.88	74.51	15.00
26	TRP		8.98	45.32	76.82	15.00
26	TRP		9.18	46.27	75.80	15.00
27	ALA	= -	6.63	42.32	75.39	15.00
27	ALA		5.70	43.30	75.95	15.00
27	ALA		4.40	42.61	76.34	15.00
27			3.92	42.75	77.47	15.00
27			5.43	44.39	74.93	15.00
28	PHE		3.83	41.86	75.40	15.00
28	PHE		2.60	41.13	75.63	15.00
	PHE		2.79	40.08	76.73	15.00
	PHE		2.00	40.02	77.66	15.00
28	PHE		2.10	40.48	74.33	15.00
28	PHE		1.41	41.44	73.39	15.00
28	PHE	CD1	2.14	42.29	72.57	15.00
28	PHE	CD2	0.02	41.51	73.33	15.00

28 PHE CE1	1.51	43.18	71 72	15 44
28 PHE CE2	-0.62		_	
28 PHE CZ	0.12			
29 SER N	3.85		·	
29 SER CA	4.15			15.00
29 SER C	4.33	38.25	· -	
29 SER 0				
29 SER CB	3.95			
29 SER OG	5.43		77.25	
30 SER N	5.78		78.20	
30 SER CA	4.96		- : • •	
30 SER CA	5.18	40.71		15.00
30 SER C	3.84	41.16		15.00
	3.48	40.84	82.03	15.00
	6.10	41.91	80.10	15.00
30 SER OG	7.39	41.48	79.72	15.00
31 VAL N	3.08	41.87	80.07	15.00
31 VAL CA	1.75	42.34	80.48	15.00
31 VAL C	0.85	41.18	80.88	15.00
31 VAL O	0.03	41.31	81.77	15.00
31 VAL CB	1.09	43.16	79.36	15.00
31 VAL CG1	-0.41	43.10	79.47	15.00
31 VAL CG2	1.57	44.60	79.43	15.00
32 GLY N	1.05	40.04	80.24	15.00
32 GLY CA	0.26	38.85	80.51	15.00
32 GLY C	0.56	38.20	81.83	15.00
32 GLY O	-0.24	37.41		15.00
33 ALA N	1.74	38.46	82.39	15.00
33 ALA CA	2.10	37.90	83.69	15.00
33 ALA C	1.61	38.88	84.75	15.00
33 ALA O	1.01	38.49	85.75	15.00
33 ALA CB	3.59	37.68	83.80	15.00
34 LEU N	1.79	40.17	84.49	15.00
34 LEU CA	1.35	41.21	85.40	15.00
34 LEU C	-0.15	41.08	85.64	15.00
34 LEU O	-0.65	41.47	86.69	15.00
34 LEU CB	1.64		84.83	
34 LEU CG	3.09	42.93	84.57	15.00
34 LEU CD1	3.08	44.29		15.00
34 LEU CD2	3.84	43.15	83.81	15.00
35 GLU N	_		85.87	15.00
35 GLU CA		40.5€	84.67	15.00
35 GLU C	-2.52	40.39	84.82	15.00
35 GLU O		39.14	85.64	15.00
35 GLU CB	-3.66 -2.98	39.11	86.37	15.00
-5 <b>-50 -5</b>	-2.38	40.28	83.45	15.00

35 GLU CG	-2.84	41.50	82.57	15.00
35 GLU CD	-3.34	41.23	81.17	15.00
35 GLU OE1	-3.19	40.08	80.70	15.00
35 GLU OE2	-3.87	42.16	80.54	15.00
36 GLY N	-1.84	38.10	85.50	15.00
36 GLY CA	-2.08	36.89	86.26	15.00
36 GLY C	-1.99	37.16	87.76	15.00
36 GLY O	-2.78	36.62	88.55	15.00
37 GLN N	-1.03	37.99	88.17	15.00
37 GLN CA	-0.86	38.31	89.57	15.00
37 GLN C	-1.88	39.32	90.06	15.00
37 GLN 0	-2.40	39.18	91.17	15.00
37 GLN CB	0.55	38.83	89.86	15.00
37 GLN CG	1.61	37.74	90.00	15.00
37 GLN CD	1.14	36.57	90.83	15.00
37 GLN OE1	0.97	36.67	92.05	15.00
37 GLN NE2	0.93	35.43	90.17	15.00
38 LEU N	-2.17	40.32	89.23	15.00
38 LEU CA	-3.14	41.34	89.61	15.00
38 LEU C	-4.49	40.73	90.02	15.00
38 LEU O	-5.12	41.20	90.95	15.00
38 LEU CB	-3.34	42.35	88.48	15.00
38 LEU CG	-4.22	43.54	88.86	15.00
38 LEU CD1	-3.64	44.20	90.08	15.00
38 LEU CD2	-4.33	44.53	87.71	15.00
39 LYS N	-4.92	39.70	89.30	15.00
39 LYS CA	-6.18	39.02	89.60	15.00
39 LYS C	-6.00	38.22	90.90	15.00
39 LYS O	-6.92	38.19	91.73	15.00
39 LYS CB	-6.58	38.09	88.45	15.00
39 LYS CG	-7.57	36.99	88.79	15.00
39 LYS CD	-8.95	37.51	89.13	15.00
39 LYS CE	-9.89	36.35	89.41	15.00
39 LYS NZ	-11.25	36.77	89.82	15.00
40 LYS N	-4.83	37.60	91.07	15.00
40 LYS CA	-4.57	36.82	92.28	15.00
40 LYS C	-4.64	37.74	93.49	15.00
40 LYS O	-5.25	37.40	94.51	15.00
40 LYS CB	-3.20	36.16	92.21	15.00
40 LYS CG	-3.09	34.85	92.98	15.00
40 LYS CD	-1.63	34.41	93.04	15.00
40 LYS CE	-1.44	33.15	93.86	15.00
40 LYS NZ	0.00	32.80	94.08	15.00
41 LYS N	-4.06	38.93	93.36	15.00

41	LYS CA	-4.06	39.93	94.43	15.00
41	LYS C	-5.40	40.66	94.59	15.00
41	LYS O	-6.17	40.38	95.49	15.00
41	LYS CB	-2.93	40.95	94.23	15.00
41	LYS CG	-1.55	40.33	94.11	15.00
41	LYS CD	-1.34	39.26	95.18	15.00
41	LYS CE	-0.06	38.46	94.95	15.00
41	LYS NZ	-0.04	37.19	95.74	15.00
42	THR N	-5.69	41.58	93.67	15.00
42	THR CA	-6.91	42.36	93.75	15.00
42	THR C	-8.23	41.65	93.40	15.00
42	THR O	-9.28	42.28	93.37	15.00
42	THR CB	-6.77	43.64	92.91	15.00
42	THR OG1	-6.89	43.34	91.51	15.00
42	THR CG2	-5.40	44.26	93.16	15.00
43	GLY N	-8.17	40.35	93.10	15.00
43	GLY CA	-9.38	39.62	92.78	15.00
43	GLY C	-10.09	39.89	91.44	15.00
43	GLY O	-10.94	39.09	91.03	15.00
44		-9.82	41.02	90.79	15.00
44		-10.45	41.31	89.50	15.00
44		-9.41	41.72	88.45	15.00
44		-8.48	42.49	88.73	15.00
44		-11.57	42.34	89.64	15.00
44		-11.20	43.55	90.45	15.00
44	LYS CD	-12.43	44.28	90.95	15.00
44	LYS CE	-12.02	45.39	91.93	15.00
44	LYS NZ	-11.26	44.88	93.11	15.00
45	LEU N	-9.60	41.19	87.25	15.00
45	LEU CA	-8.69	41.41	86.13	15.00
45	LEU C	-8.92	42.66	85.26	15.00
45	LEU O	-10.04	43.15	85.12	15.00
45	LEU CB	-8.71	40.16	85.25	15.00
	LEU CG	-7.54	39.90	84.33	15.00
45		-6.25	39.87	85.12	15.00
45		-7.75	38.59	83.63	15.00
	LEU N	-7.83	43.15	84.69	15.00
46	LEU CA	-7.84	44.31	83.80	15.00
46	LEU C	-6.81	44.09	82.69	15.00
46	LEU O	-5.76	43.51	82.93	15.00
	LEU CB	-7.44	45.58	84.55	15.00
	LEU CG	-8.49	46.47	85.20	15.00
46	LEU CD1	-7.96	47.90	85.21	15.00
46	LEU CD2	-9.77	46.43	84.42	15.00

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47 ASN N	-7.11	44.56	81.49	15.00
47 ASN CA	-6.17	44.43	80.38	15.00
47 ASN C	-5.12	45.50	80.56	15.00
47 ASN O	-5.45	46.67	80.62	15.00
47 ASN CB	-6.88	44.66	79.03	15.00
47 ASN CG	-7.67	43.45	78.57	15.00
47 ASN OD1	-7.13	42.37	78.38	15.00
47 ASN ND2	-8.97	43.65		15.00
48 LEU N	-3.86	45.10	80.70	15.00
48 LEU CA	-2.77	46.07	80.88	15.00
48 LEU C	-2.15	46.34	79.52	15.00
48 LEU O	-2.28	45.53	78.61	15.00
48 LEU CB	-1.74	45.56		15.00
48 LEU CG	-2.26	45.06	83.27	15.00
48 LEU CD1	-1.09	44.75	84.18	15.00
48 LEU CD2	-3.18	46.08	83.91	15.00
49 SER N	-1.46	47.47	79.36	15.00
49 SER CA	-0.91	47.82	78.06	15.00
49 SER C	0.51	47.42	77.67	15.00
49 SER O	1.48	47.79	78.33	15.00
49 SER CB	-1.10	49.32	77.77	15.00
49 SER OG	-0.19	50.13	78.51	15.00
50 PRO N	0.64	46.66	76.56	15.00
50 PRO CA	1.94	46.22	76.04	15.00
50 PRO C	2.59	47.44	75.41	15.00
50 PRO O	3.80	47.52	75.25	15.00
50 PRO CB	1.54	45.21	74.97	15.00
50 PRO CG	0.21	44.71	75.44	15.00
50 PRO CD	-0.45	45.98	75.85	15.00
51 GLN N	1.74	48.40	75.03	15.00
51 GLN CA	2.18	49.65	74.43	15.00
51 GLN C	3.01	50.45	75.42	15.00
51 GLN 0	4.12	50.87	75.11	15.00
51 GLN CB	0.97	50.49	74.00	15.00
51 GLN CG	1.33	51.66	73.08	15.00
51 GLN CD	1.77	51.20	71.71	15.00
51 GLN OE1	0.96	50.74	70.92	15.00
51 GLN NE2	3.05	51.26	71.44	15.00
52 ASN N	2.47	50.66	76.62	15.00
52 ASN CA	3.18	51.42	77.65	15.00
52 ASN C	4.59	50.85	77.83	15.00
52 ASN 0	5.56	51.59	77.97	15.00
52 ASN CB	2.42	51.35	78.97	15.00
52 ASN CG	3.09	52.12	80.09	15.00

52 ASN OD1	2.74	51.95	81.25	15.00
52 ASN ND2	4.04	52.99		
53 LEU N	4.68	49.53	77.81	15.00
53 LEU CA	5.95	48.86	77.94	15.00
53 LEU C	6.82	49.21	76.72	15.00
53 LEU O	7.87	49.84	76.88	
53 LEU CB	5.75	47.35	78.08	15.00
53 LEU CG	5.11	46.95	79.41	15.00
53 LEU CD1	4.91	45.45	79.49	15.00
53 LEU CD2	6.00	47.41	80.54	15.00
54 VAL N	6.33	48.87		15.00
54 VAL CA	7.03	49.15		15.00
54 VAL C	7.63	50.55	74.20	15.00
54 VAL O	8.85	50.72	74.04	15.00
54 VAL CB	6.06	49.01	73.03	15.00
54 VAL CG1	6.70	49.55	71.75	15.00
54 VAL CG2	5.67	47.56	72.83	15.00
55 ASP N	6.76	51.55	74.37	15.00
55 ASP CA	7.12	52.96	74.31	15.00
55 ASP C	8.05	53.45	75.40	15.00
55 ASP 0	8.84	54.37	75.19	15.00
55 ASP CB	5.85	53.84	74.36	15.00
55 ASP CG	4.87	53.56	73.22	15.00
55 ASP OD1	5.23	52.86	72.24	15.00
55 ASP OD2	3.72	54.05	73.29	15.00
56 CYS N	7.97	52.82	76.56	15.00
56 CYS CA	8.76	53.27	77.70	15.00
56 CYS C	9.97	52.47	78.16	15.00
56 CYS O	10.98	53.06	78.54	15.00
56 CYS CB	7.81	53.52	78.84	15.00
56 CYS SG	6.36	54.45	78.28	15.00
57 VAL N	9.87	51.15	78.17	15.00
57 VAL CA	10.98	50.31	78.60	15.00
57 VAL C	12.16	50.44	77.62	15.00
57 VAL O	12.50	49.50	76.90	15.00
57 VAL CB	10.54	48.83	78.71	15.00
57 VAL CG1	11.66	47.98	79.32	15.00
57 VAL CG2	9.26	48.72	79.52	15.00
58 SER N	12.85	51.57	77. <b>67</b>	15.00
58 SER CA	13.98	51.87	76.80	15.00
58 SER C	15.15	50.91	76.84	15.00
58 SER O	16.13	51.10	76.13	15.00
58 SER CB	14.48	53.27	77.09	15.00
58 SER OG	14.95	53.34	78.42	15.00

59	GLU	N	15.09	49.92	77.71	15.00
59	GLU	CA	16.15	48.93	77.82	15.00
59	GLU	С	15.93	47.86	76.73	15.00
59	GLU	0	16.77	46.99	76.50	15.00
59	GLU	CB	16.14	48.30	79.22	15.00
59	GLU	CG	16.39	49.27	80.39	15.00
59	GLU	CD	15.13	49.97	80.88	15.00
59	GLU	OE1	14.38	49.38	81.70	15.00
59	GLU	OE2	14.90	51.13	80.46	15.00
60	ASN	N	14.77	47.91	76.10	15.00
60	ASN	CA	14.43	46.98	75.04	15.00
60	ASN	C	14.26	47.73	73.71	15.00
60	ASN	0	13.99	48.94	73.69	15.00
60	ASN	CB	13.13	46.26	75.38	15.00
60	ASN	CG	13.27	45.35	76.57	15.00
60	ASN	OD1	12.35	45.22	77.38	15.00
60	ASN	ND2	14.41	44.68	76.68	15.00
61	ASP	N	14.39	47.02	72.61	15.00
61	ASP	CA	14.25	47.65	71.31	15.00
61	ASP	С	12.82	48.03	70.94	15.00
61	ASP	0	12.60	48.65	69.91	15.00
61	ASP	СВ	14.84	46.75	70.23	15.00
61	ASP	CG	15.93	47.43	69.44	15.00
61	ASP	OD1	15.87	48.66	69.28	15.00
61	ASP	OD2	16.85	46.72	68.98	15.00
62	GLY	N	11.84	47.67	71.77	15.00
62	GLY	CA	10.46	47.99	71.45	15.00
62	GLY	С	9.91	47.03	70.41	15.00
62	GLY	0	9.71	45.85	70.70	15.00
	CYS	N	9.68	47.50	69.20	15.00
63	CYS		9.19	46.60	68.16	15.00
63	CYS		10.33	45.71	67.70	15.00
63	CYS	0	10.09	44.70	67.04	15.00
63	CYS	CB	8.59	47.36	66.99	15.00
63	CYS	SG	6.94	48.02	67.38	15.00
	GLY		11.56	46.10	68.03	15.00
64	GLY	CA	12.72	45.32	67.68	15.00
64	GLY	C	12.90	44.16	68.64	15.00
64	GLY	0	13.65	43.21	68.37	15.00
65	GLY	N	12.20	44.23	69.78	15.00
65	GLY	CA	12.28	43.17	70.77	15.00
65	GLY	C	12.86	43.58	72.11	15.00
65	$GL_iY$	0	13.34	44.70		15.00
66	GLY	N	12.81	42.65	73.05	15.00

66	GLY	CA	13.34	42.92	74.37	15.00
66	GLY	С	13.08	41.77	75.32	15.00
66	GLY	0	12.42	40.80	74.94	15.00
67	TYR	N	13.57	41.90	76.55	15.00
67	TYR	CA	13.40	40.87	77.56	15.00
67	TYR	С	12.23	41.18	78.48	15.00
67	TYR	0	11.93	42.34	78.75	15.00
67	TYR	CB	14.68	40.76	78.39	15.00
67	TYR	CG	15.91	40.56	77.55	15.00
67	TYR	CD1	16.20	39.32	76.98	15.00
67	TYR	CD2	16.77	41.61	77.29	15.00
67	TYR	CE1	17.33	39.13	76.18	15.00
67	TYR	CE2	17.90	41.44	76.49	15.00
67	TYR	CZ	18.18	40.20	75.94	15.00
67	TYR	OH	19.27	40.05	75.12	15.00
68	MET	N	11.57	40.13	78.99	15.00
68	MET	CA	10.45	40.33	79.90	15.00
68	MET	С	10.90	40.92	81.23	15.00
68	MET	0	10.21	41.72	81.83	15.00
68	MET	CB	9.72	39.00	80.18	15.00
68	MET	CG	8.97	38.38	79.00	15.00
	MET		9.97	37.36	77.92	15.00
	MET	CE	11.01	36.45	79.11	15.00
69	THR	N	12.09	40.53	81.68	15.00
69			12.62	41.03	82.94	15.00
69	THR	С	12.76	42.55	82.93	15.00
69	THR		12.33	43.21	83.87	15.00
69	THR		13.98	40.37	83.29	15.00
69		OG1	14.96	40.72	82.30	15.00
69		CG2	13.84	38.85	83.35	15.00
70			13.33	43.09	81.85	15.00
	ASN		13.53	44.54	81.73	15.00
			12.24	45.34	81.80	15.00
70		0	12.25	46.51	82.16	15.00
	ASN		14.28	44.87	80.45	15.00
70			15.72	44.45	80.52	15.00
	asn		16.11	43.68	81.39	15.00
	ASN		16.54	44.97	79.61	15.00
	ALA		11.14	44.68	81.45	15.00
71			9.81	45.28	81.50	15.00
	ALA		9.27	45.18	82.93	15.00
	ALA		8.72	46.14	83.46	15.00
	ALA		8.83	44.57	80.51	15.00
72	PHE	N	9.46	44.02	83.55	15.00

72	PHE	CA	9.02	43.79	84.93	15.00
72	PHE	C	9.63	44.86	85.81	15.00
72	PHE	0	8.94	45.51	86.59	15.00
72	PHE	CB	9.49	42.43	85.44	15.00
72	PHE	CG	8.73	41.28	84.88	15.00
72	PHE	CD1	7.40	41.42	84.52	15.00
72	PHE	CD2	9.34	40.04	84.73	15.00
72	PHE	CE1	6.67	40.35	84.03	15.00
72	PHE	CE2	8.62	38.96	84.24	15.00
72	PHE	CZ	7.29	39.12	83.89	15.00
73	GLN	N	10.93	45.05	85.64	15.00
73	GLN	CA	11.67	46.04	86.41	15.00
73	GLN	C	11.17	47.48	86.11	15.00
73	GLN	0	11.29	48.37	86.94	15.00
73	GLN	CB	13.16	45.86	86.13	15.00
73	GLN	CG	14.11	46.75	86.94	15.00
73	GLN	CD	14.52	46.20	88.32	15.00
73	GLN	OE1	15.45	46.73	88.93	15.00
73	GLN	NE2	13.85	45.16	88.80	15.00
74	TYR	N	10.54	47.68	84.96	15.00
74	TYR	CA	10.04	49.00	84.58	15.00
74	TYR	С	8.81	49.41	85.38	15.00
74	TYR	0	8.69	50.56	85.81	15.00
74	TYR	CB	9.72	49.05	83.08	15.00
74	TYR	CG	8.90	50.26	82.67	15.00
74	TYR		9.48	51.52	82.60	15.00
74	TYR	CD2	7.54	50.14	82.44	15.00
74			8.71	52.63	82.31	15.00
74	TYR	CE2	6.77	51.25	82.15	15.00
74	TYR	CZ	7.36	52.49	82.09	15.00
	TYR		6.58	53 . 59	81.84	15.00
	VAL		7.87	49.48	85.5 <b>4</b>	15.00
	VAL		6.65	48.74	86.31	15.00
	VAL	С	7.07	49.05	87.76	15.00
	VAL		6.41	49.80	88.47	15.00
75	VAL	CB	5.73	47.48	86.38	15.00
	VAL		4.32	47.87	86.73	15.00
75	VAL	CG2	5.77	46.72	85.07	15.00
	GLN	N	8.19	43.44	88.18	15.00
76	GLN	CA.	8.71	48.62	89.52	15.00
76		С	9.26	50.02	89.71	15.00
		0	8.62	50.86	90.31	15.00
	GLN		9.78	47.57	89.79	15.00
76	GLN	CG	10.35	47.60	91.20	15.00

76 GLN CD	11.53	46.64	91.36	15.00
76 GLN OE1	12.17	46.25	90.38	15.00
76 GLN NE2	11.80	46.25	92.59	15.00
77 LYS N	10.43	50.30	89.15	15.00
77 LYS CA	11.04	51.62	89.32	15.00
77 LYS C	10.24	52.83	88.85	15.00
77 LYS 0	10.34	53.90	89.44	15.00
77 LYS CB	12.44	51.64	88.71	15.00
77 LYS CG	12.52	51.06	87.30	15.00
77 LYS CD	13.96	50.69	86.97	15.00
77 LYS CE	14.06	49.75	85.79	15.00
77 LYS NZ	15.39	49.07	85.82	15.00
78 ASN N	9.48	52.69	87.77	15.00
78 ASN CA	8.67	53.81	87.31	15.00
78 ASN C	7.51	53.96	88.28	15.00
78 ASN 0	6.94	55.03	88.41	15.00
78 ASN CB	8.12	53.56	85.90	15.00
78 ASN CG	7.17	54.66	85.44	15.00
78 ASN OD1	7.61	55.76	85.09	15.00
78 ASN ND2	5.88	54.37	85.44	15.00
79 ARG N	7.19	52.86	88.95	15.00
79 ARG CA	6.11	52.78	89.94	15.00
79 ARG C	4.71	52.88	89.33	15.00
79 ARG O	4.05	53.92	89.37	15.00
79 ARG CB	6.31	53.81	91.08	15.00
79 ARG CG	7.67	53.69	91.80	15.00
79 ARG CD	7.80	54.56	93.06	15.00
79 ARG NE	6.84	54.23	94.12	15.00
79 ARG CZ	6.60	53.00	94.59	15.00
79 ARG NH1	7.24	51.94	94.09	15.00
79 ARG NH2	5.73	52.83	95.58	15.00
80 GLY N	4.27	51.76	88.76	15.00
80 GLY CA	2.95	51.68	88.15	15.00
80 GLY C	2.98	51.54	86.64	15.00
80. GLY O	3.87	52.07	85.97	15.00
81 ILE N	1.98	50.85	86.10	15.00
81 ILE CA	1.88	50.64	84.66	15.00
81 ILE C	0.45	50.97	84.23	15.00
81 ILE O	-0.49	50.68	84.96	15.00
81 ILE CB	2.19	49.17	84.30	15.00
81 ILE CG1	2.27	48.99	82.78	15.00
81 ILE CG2	1.14	48.24	84.90	15.00
81 ILE CD1	2.64	47.59	82.36	15.00
82 ASP N	0.29	51.58	83.05	15.00

82	ASP	CA	-1.04	51.93	82.56	15.00
82	ASP	С	-1.89	50.76	82.07	15.00
82	ASP	0	-1.38	49.69	81.71	15.00
82	ASP	CB	-0.93	52.93	81.42	15.00
82	ASP	CG	-0.47	54.29	81.87	15.00
82	ASP	OD1	-0.66	54.63	83.06	15.00
82	ASP	OD2	0.08	55.02	81.02	15.00
83	SER	N	-3.20	50.96	82.05	15.00
83	SER	CA	-4.12	49.95	81.56	15.00
83	SER	С	-4.32	50.20	80.07	15.00
83	SER	0	-4.22	51.34	79.61	15.00
83	SER	CB	-5.46	50.02	82.32	15.00
83	SER	OG	-6.06	51.30	82.22	15.00
84	GLU	N	-4.61	49.14	79.32	15.00
84	GLU	CA	-4.81	49.22	77.86	15.00
84	GLU	С	-5.66	50.40	77.38	15.00
84	GLU	0	-5.20	51.19	76.57	15.00
84	GLU	CB	-5.39	47.91	77.33	15.00
	GLU		-5.73	47.89	75.85	15.00
84	GLU		-4.51	47.97	74.94	15.00
84		OE1	-3.51	47.25	75.17	15.00
84		OE2	-4.57	48.74	73.96	15.00
	ASP	N	-6.87	50.53	77.90	15.00
85	ASP	CA	-7.72	51.61	77.45	15.00
85	ASP	С	-7.12	52.99	77.70	15.00
	ASP		-7.49	53.97	77.03	15.00
	ASP		-9.10	51.52	78.07	15.00
	ASP		-10.07	52.52	77.46	15.00
	ASP		-9.97	52.79	76.24	15.00
	ASP		-10.93	53.06	78.21	15.00
	ALA		-6.18	53.08	78.64	15.00
	ALA		-5.54	54.36	78.93	15.00
	ALA		-4.30	54.57	78.04	15.00
	ALA		-3.90	55.70	77.78	15.00
	ALA		-5.16	54.45		15.00
	TYR		-3.74	53.48	77.53	15.00
87		CA	-2.56	53.54	76.67	15.00
87			-2.69	52.41		15.00
87			-1.98	51.39		15.00
87		CB	-1.32	53.33	77.55	15.00
87		CG	-0.03	53.89		15.00
87		CD1	0.20	53.99		15.00
87		CD2		54.29		15.00
87	TYR	CE1	1.41	54.47	75.16	15.00

87	TYR	CE2	2.20	54.77	77.41	15.00
87	TYR	CZ	2.41	54.86	76.05	15.00
87	TYR	OH	3.61	55.32	75.57	15.00
88	PRO	N	-3.60	52.58	74.65	15.00
88	PRO	CA	-3.93	51.66	73.55	15.00
88	PRO	С	-2.80	51.22	72.61	15.00
88	PRO	0	-2.08	52.05	72.06	15.00
88	PRO		-5.01	52.43	72.79	15.00
88	PRO	CG	-5.64	53.28	73.84	15.00
88	PRO	CD	-4.42	53.80	74.54	15.00
89	TYR	N	-2.74	49.92	72.34	15.00
89	TYR	CA	-1.71	49.35	71.48	15.00
89	TYR	С	-1.83	49.79	70.01	15.00
89	TYR	0	-2.46	49.11	69.21	15.00
89	TYR	CB	-1.74	47.82	71.56	15.00
89	TYR	CG	-0.54	47.18	70.92	15.00
89	TYR	CD1	0.75	47.59	71.30	15.00
89	TYR	CD2	-0.66	46.24	69.91	15.00
89	TYR	CE1	1.88	47.07	70.69	15.00
89	TYR	CE2	0.47	45.70	69.28	15.00
89	TYR	CZ	1.74	46.13	69.68	15.00
89	TYR	OH	2.87	45.65	69.09	15.00
90	VAL	N	-1.19	50.88	69.65	15.00
90	VAL	CA	-1.22	51.40	68.29	15.00
90	VAL	С	-0.55	50.51	67.23	15.00
90	VAL	0	-0.83	50.63	66.03	15.00-
90	VAL	CB	-0.63	52.83	68.21	15.00
90	VAL	CG1	-1.26	53.71	69.28	15.00
90	VAL	CG2	0.88	52.80	68.33	15.00
91	GLY	N	0.37	49.65	67.64	15.00
91	GLY	CA	1.01	48.77	66.67	15.00
91	GLY	С	2.38	49.17	66.15	15.00
91	GLY	0	2.89	48.57	65.20	15.00
92	GLN	N	3.00	50.17	66.77	15.00
92	GLN	CA	4.32	50.61		15.00
92	GLN	C	4.96	51.34	67.52	15.00
92	GLN	0	4.33	51.51	68.57	15.00
92	GLN	CB	4.21	51.53	65.13	15.00
93	GLN	CG	3.46	52.85	65.38	15.00
92	GLN	CD	3.28	53.70	64.12	15.00
92	GLN	OE1	3.93	54.73	63.94	15.00
92	GLN	NE2	2.39	53.25	63.25	15.00
93	GLU	J N	6.19	51.79	67.34	15.00
93	GLU	J CA	6.90	52.49	68.39	15.00

93 GLU C	6.60	53.99	68.38	15.00
93 GLU 0	6.84	54.67	67.38	15.00
93 GLU CB	8.41	52.27	68.23	15.00
93 GLU CG	8.80	50.81	68.17	15.00
93 GLU CD	10.30	50.60	68.23	15.00
93 GLU OE1	10.85	50.53	69.35	15.00
93 GLU OE2	10.93	50.50	67.15	15.00
94 GLU N	6.08	54.49	69.49	15.00
94 GLU CA	5.76	55.90	69.63	15.00
94 GLU C	6.54	56.47	70.80	15.00
94 GLU 0	7.40	55.80	71.37	15.00
94 GLU CB	4.27	56.10	69.86	15.00
94 GLU CG	3.45	56.13	68.58	15.00
94 GLU CD	1.96	56.28	68.85	15.00
94 GLU OE1	1.48	55.64	69.81	15.00
94 GLU OE2	1.27	57.03	68.11	15.00
95 SER N	6.22	57.71	71.14	15.00
95 SER CA	6.84	58.41	72.26	15.00
95 SER C	6.16	57.91	73.52	15.00
95 SER O	4.92	57.82	73.58	15.00
95 SER CB	6.63	59.93	72.12	15.00
95 SER OG	5.25	60.23	71.89	15.00
96 CYS N	6.96	57.55	74.51	15.00
96 CYS CA	6.44	57.04	75.77	15.00
96 CYS C	5.44	58.02	76.39	15.00
96 CYS 0	5.84	59.03	76.95	15.00
96 CYS CB	7.59	56.77	76.74	15.00
96 CYS SG	7.00	56.38	78.40	15.00
97 MET N	4.15	57.73	76.21	15.00
97 MET CA	3.06	58.55	76.74	15.00
97 MET C	2.40	58.00	78.01	15.00
97 MET O	1.16	57.90	78.06	15.00 15.00
97 MET CB	1.97	58.73	75.69 74.45	15.00
97 MET CG	2.36	59.52	73.09	15.00
97 MET SD	1.29	58.98	73.09	15.00
97 MET CE	-0.36	59. <b>43</b> 57.65	79.01	15.00
98 TYR N	3.20		80.26	15.00
98 TYR CA	2.67	57.13	81.06	15.00
98 TYR C	1.93	58.21	81.38	15.00
98 TYR O	2.48	59.26 56.51		15.00
98 TYR CB	3.78	56.04	82.48	15.00
93 TYR CG	3.32 3.24	56.92		15.00
98 TYR CD1	2.95			
98 TYR CD2	2.33	24.14	02.00	

98 TYR CE1	2.81	56.49	84.80	15.00
98 TYR CE2	2.52	54.29	83.95	15.00
98 TYR CZ	2.45	55.18	84.99	15.00
98 TYR OH	2.02	54.77	86.22	15.00
99 ASN N	0.69	57.92	81.43	15.00
99 ASN CA	-0.14	58.84	82.20	15.00
99 ASN C	-0.54	58.20	83.54	15.00
99 ASN 0	-1.31	57.23	83.58	15.00
99 ASN CB	-1.37	59.22	81.36	15.00
99 ASN CG	-2.38	60.08	82.13	15.00
99 ASN OD1	-2.01	60.89	82.99	15.00
99 ASN ND2	-3.66	59.90	81.80	15.00
100 PRO N	-0.02	58.74	84.65	15.00
100 PRO CA	-0.30	58.24	86.00	15.00
100 PRO C	-1.77	58.23	86.33	15.00
100 PRO O	-2.22	57.46	87.19	15.00
100 PRO CB	0.45	59.22	86.90	15.00
100 PRO CG	1.58	59.70	86.02	15.00
100 PRO CD	0.86	59.92	84.71	15.00
101 THR N	-2.53	59.08	85.65	15.00
101 THR CA	-3.96	59.13	85.89	15.00
101 THR C	-4.56	57.81	85.43	15.00
101 THR O	-5.64	57.43	85.89	15.00
101 THR CB	-4.63	60.25	85.10	15.00
101 THR OG1	-3.86	61.45	85.21	15.00
101 THR CG2	-6.03	60.52	85.66	15.00
102 GLY N	-3.86	57.13	84.52	15.00
102 GLY CA	-4.33	55.86	84.01	15.00
102 GLY C	-3.55	54.62	84.42	15.00
102 GLY O	-3.52	53.65	83.67	15.00
103 LYS N	-2.90	54.62	85.59	15.00
103 LYS CA	-2.15	53.44	86.02	15.00
103 LYS C	-3.12	52.35	86.50	15.00
103 LYS O	-4.26	52.65	86.86	15.00
103 LYS CB	-1.13	53.83	87.11	15.00
103 LYS CG	-1.51	53.50	88.56	15.00
103 LYS CD	-0.97	52.13	88.98	15.00
103 LYS CE	-1.39	51.77	90.40	15.00
103 LYS NZ	-1.11	50.35	90.75	15.00
104 ALA N	-2.68	51.10	86.52	15.00
104 ALA CA	-3.55	50.01	86.96	15.00
104 ALA C	-2.86	48.86	87.68	15.00
104 ALA O	-3.52	47.92	88.12	15.00
104 ALA CB	-4.37	49.49	85.79	15.00

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105	ALA	N	-1.55	48.92	87.82	15.00
105	ALA	CA	-0.83	47.87	88.50	15.00
105	ALA	С	0.53	48.36	88.97	15.00
105	ALA	0	1.06	49.35	88.44	15.00
105	ALA	CB	-0.67	46.68	87.59	15.00
106	LYS	N	1.07	47.69	90.00	15.00
106	LYS	CA	2.39	48.03	90.56	15.00
106	LYS	C	3.24	46.76	90.62	15.00
106	LYS	0	2.73	45.65	90.44	15.00
106	LYS	CB	2.24	48.66	91.95	15.00
106	LYS	CG	1.78	50.12	91.94	15.00
106	LYS	CD	1.32	50.56	93.31	15.00
106	LYS	CE	2.44	50.52	94.34	15.00
106	LYS	NZ	3.48	51.56	94.06	15.00
107	CYS	N	4.54	46.92	90.90	15.00
	CYS		5.42	45.76	90.90	15.00
	CYS		6.51	45.83	91.97	15.00
	CYS		7.16	46.86	92.14	15.00
	CYS		6.07	45.65	89.52	15.00
	CYS		6.62	44.03	89.03	15.00
	ARG		6.70	44.73	92.69	15.00
	ARG			44.62	93.74	15.00
	ARG		8.93	43.86	93.19	15.00 15.00
	ARG		9.61	43.14	93.93	15.00
	ARG		7.16	43.87	94.96	15.00
	ARG		6.08	44.58	95.79 96.74	15.00
	ARG		5.41	43.58	96.7 <b>4</b> 97.25	15.00
	ARG		6.38	42.60	97.25	15.00
	ARG		6.16	41.29	97.08	15.00
		NH1	4.99	40.74 40.52	97.80	15.00
	ARG		7.16 9.21	44.02	91.90	15.00
	GLY GLY		10.34	43.34	91.30	15.00
	GLY		9.92	42.13	90.50	15.00
			8.77	42.01	90.10	15.00
	GLY TYR		10.86	41.22	90.29	15.00
110		CA	10.59	39.99	89.54	15.00
	TYR		11.44	38.82	90.02	15.00
	TYR		12.41	38.99	90.75	15.00
	TYP		10.85	40.21	88.05	15.00
	TYF		12.30	40.42	87.70	15.00
		CD1	13.13	39.33	87.46	15.00
		CD2	12.82	41.70	87.58	15.00
		R CE1	14.46	39.52	87.10	15.00
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110 TYR CE2	14.14	41.89	87.22	15.00
110 TYR CZ	14.95	40.80	86.98	15.00
110 TYR OH	16.25	40.99	86.61	15.00
111 ARG N	11.13	37.63	89.52	15.00
111 ARG CA	11.87	36.45	89.91	15.00
111 ARG C	11.88	35.46	88.75	15.00
111 ARG 0	10.90	35.34	88.03	15.00
111 ARG CB	11.23	35.85	91.16	15.00
111 ARG CG	12.14	34.90	91.88	15.00
111 ARG CD	11.70	34.66	93.31	15.00
111 ARG NE	12.85	34.20	94.09	15.00
111 ARG CZ	13.87	34.98	94.44	15.00
111 ARG NH1	13.87	36.28	94.12	15.00
111 ARG NH2	14.89	34.48	95.13	15.00
112 GLU N	13.00	34.76	88.58	15.00
112 GLU CA	13.15	33.79	87.50	15.00
112 GLU C	13.25	32.35	87.94	15.00
112 GLU 0	13.55	32.06	89.11	15.00
112 GLU CB	14.38	34.11	86.66	15.00
112 GLU CG	14.35	35.49	86.11	15.00
112 GLU CD	15.46	35.72	85.14	15.00
112 GLU OE1	15.30	35.30	83.97	15.00
112 GLU OE2	16.49	36.31	85.56	15.00
113 ILE N	12.98	31.46	86.99	15.00
113 ILE CA	13.02	30.03	87.20	15.00
113 ILE C	14.46	29.65	86.91	15.00
113 ILE 0	15.13	30.34	86.15	15.00
113 ILE CB	12.04	29.30	86.23	15.00
113 ILE CG1	10.60	29.50	86.70	15.00
113 ILE CG2	12.36	27.81	86.13	15.00
113 ILE CD1	10.10	30.93	86.65	15.00
114 PRO N	15.01	28.64	87.60	15.00
114 PRO CA	16.40	28.30	87.29	15.00
114 PRO C	16.50	27.74	85.87	15.00
114 PRO 0	15.74	26.85	85.48	15.00
114 PRO CB	16.77	27.29	88.39	15.00
114 PRO CG	15.46	26.66	88.76	15.00
114 PRO CD	14.52	27.85	88.75	15.00
115 GLU N	17.41	28.32	85.09	15.00
115 GLU CA	17.61	27.97	83.68	15.00
115 GLU C	17.55	26.49	83.35	15.00
115 GLU 0	18.36	25.70	83.84	15.00
115 GLU CB	18.93	28.56	83.15	15.00
115 GLU CG	18.92	30.09	82.86	15.00

115	GLU	CD	20.11	30.56	81.97	15.00
115	GLU	OE1	20.75	29.70	81.31	15.00
115	GLU	OE2	20.39	31.79	81.92	15.00
116	GLY	N	16.55	26.11	82.56	15.00
116	GLY	CA	16.41	24.74	82.12	15.00
116	GLY	С	15.76	23.78	83.08	15.00
116	GLY	0	15.81	22.56	82.88	15.00
117	ASN	N	15.13	24.31	84.11	15.00
117	ASN	CA	14.47	23.47	85.10	15.00
117	ASN	С	12.97	23.47	84.87	15.00
117	ASN	0	12.28	24.43	85.22	15.00
117	ASN	CB	14.77	23.97	86.51	15.00
117	ASN	CG	14.21	23.05	87.58	15.00
117	ASN	OD1	13.32	22.23	87.32	15.00
117	ASN	ND2	14.72	23.19	88.80	15.00
118	GLU	N	12.47	22.38	84.29	15.00
118	GLU	CA	11.05	22.27	84.03	15.00
118	GLU	С	10.22	22.01	85.27	15.00
118	GLU	0	9.16	22.60	85.42	15.00
118	GLU	CB	10.75	21.20	82.99	15.00
118	GLU	CG	10.96	21.66	81.56	15.00
118	GLU	CD	10.34	20.72	80.56	
118	GLU	OE1	9.11	20.82	80.33	15.00
118	GLU	OE2	11.08	19.88	80.01	15.00
119	LYS	N	10.73	21.18	86.18	15.00
119	LYS	CA	10.00	20.87	87.41	15.00
119	LYS	С	9.79	22.11	88.28	15.00
119	LYS	0	8.78	22.21	88.98	15.00
119	LYS	CB	10.68	19.76	88.21	15.00
119	LYS	CG	9.94	18.40	88.14	15.00
119	LYS	CD	10.13	17.64	86.81	15.00
119	LYS	CE	9.28	16.35	86.76	15.00
119	LYS	NZ	9.62	15.29	87.79	15.00
120	ALA	N	10.73	23.04	88.21	15.00
120	ALA	CA	10.61	24.30	88.94	15.00
120	ALA	C	9.57	25.15	88.23	15.00
120	ALA	0	8.72	25.75	88.88	15.00
120	ALA	CB	11.93	25.04	88.99	15.00
121	LEU	JN	9.62	25.19	86.89	15.00
121	LEU	J CA	8.66	25.96	86.09	15.00
121	LEU	JC	7.23	25.51	86.39	15.00
121	LET	JO	6.34	26.34	86.52	15.00
121	LET	J CB	8.94	25.82	84.58	15.00
121	LEU	J CG	7.91	26.43	83.60	15.00

121	LEU	CD1	8.09	27.93	83.52	15.00
121	LEU	CD2	8.08	25.84	82.22	15.00
122	LYS	N	7.02	24.21	86.51	15.00
122	LYS	CA	5.69	23.70	86.81	15.00
	LYS	С	5.24	24.18	88.19	15.00
122	LYS	0	4.07	24.52	88.38	15.00
122	LYS	CB	5.66	22.17	86.76	15.00
122	LYS	CG	4.31	21.58	87.17	15.00
122	LYS	CD	4.36	20.07	87.30	15.00
122	LYS	CE	5.50	19.63	88.22	15.00
122	LYS	NZ	5.63	18.14	88.27	15.00
123	ARG	N	6.16	24.19	89.15	15.00
123	ARG	CA	5.84	24.63	90.51	15.00
123	ARG	С	5.51	26.11	90.47	15.00
123	ARG	0	4.53	26.54	91.07	15.00
123	ARG	CB	6.99	24.36	91.49	15.00
123	ARG	CG	7.32	22.88	91.65	15.00
123	ARG	CD	8.14	22.58	92.89	15.00
123	ARG	NE	9.59	22.54	92.68	15.00
123	ARG	CZ	10.40	23.60	92.76	15.00
123	ARG	NH1	9.92	24.80	93.04	15.00
123	ARG	NH2	11.71	23.43	92.63	15.00
124	ALA	N	6.30	26.87	89.73	15.00
	ALA		6.09	28.30	89.58	15.00
	ALA		4.72	28.57	88.98	15.00
	ALA		3.98	29.41	89.47	15.00
	ALA		7.19	28.91	88.71	15.00
	VAL		4.37	27.84	87.92	15.00
	VAL		3.07	28.02	87.28	15.00
	VAL		1.95	27.63	88.24	15.00
	VAL		0.88	28.23	88.22	15.00
125	VAL		2.96	27.17	85.98	15.00 15.00
125		. CG1	1.52	27.07	85.51	15.00
		. CG2	3.78	27.79	84.88	15.00
	5 ALA		2.21	26.64	89.08	15.00
	5 ALA		1.22	26.16	90.04	15.00
	5 AL		1.10	26.92	91.36	15.00
	5 AL		0.03	26.95	91.97	15.00
	S ALA		1.43	24.68	90.31	15.00
12		3 N	2.20	27.51	91.82	15.00
12		G CA	2.21	28.25	93.07	15.00
12			1.97	29.74	92.86	15.00
12			1.45	30.43	93.75	
12	7 AR	G CB	3.55	28.07	93.81	15.00

127 ARG CG	3.44	27.25	95.07	15.00
127 ARG CD	3.83	25.82	94.84	15.00
127 ARG NE	5.24	25.57	95.18	15.00
127 ARG CZ	5.76	24.37	95.39	15.00
127 ARG NH1	5.03	23.27	95.29	15.00
127 ARG NH2	7.04	24.27	95.69	15.00
128 VAL N	2.38	30.23	91.69	15.00
128 VAL CA	2.22	31.64	91.36	15.00
128 VAL C	1.08	31.86	90.37	15.00
128 VAL O	0.06	32.47	90.67	15.00
128 VAL CB	3.53	32.20	90.78	15.00
128 VAL CG1	3.48	33.72	90.75	15.00
128 VAL CG2	4.72	31.71	91.58	15.00
129 GLY N	1.27	31.35	89.16	15.00
129 GLY CA	0.26	31.49	88.14	15.00
129 GLY C	0.98	31.73	86.84	15.00
129 GLY 0	2.11	31.28	86.67	15.00
130 PRO N	0.36	32.47	85.91	15.00
130 PRO CA	0.96	32.76	84.62	15.00
130 PRO C	2.37	33.32	84.72	15.00
130 PRO O	2.60	34.37	85.32	15.00
130 PRO CB	-0.03	33.76	84.02	15.00
130 PRO CG	-1.33	33.27	84.53	15.00
130 PRO CD	-1.00	33.01	85.99	15.00
131 VAL N	3.32	32.57	84.15	15.00
131 VAL CA	4.72	32.96	84.13	15.00
131 VAL C	5.09	33.29	82.67	15.00
131 VAL 0	4.48	32.77	81.73	15.00
131 VAL CB	5.63	31.83	84.73	15.00
131 VAL CG1	5.57	30.58	83.89	15.00
131 VAL CG2	7.07	32.28	84.86	15.00
132 SER N	6.03	34.23	82.49	15.00
132 SER CA	6.49	34.64	81.17	15.00
132 SER C	7.58	33.70	80.71	15.00
132 SER O	8.60	33.56	81.40	15.00
132 SER CB	7.08	36.05	81.25	15.00
132 SER OG	6.16	36.98	81.80	15.00
133 VAL N	7.39	33.03	79.58	15.00
133 VAL CA	8.39	32.10	79.03	15.00
133 VAL C	8.84	32.58	77.67	15.00
133 VAL 0	8.20	33.43	77.06	15.00
133 VAL CB	7.86	30.6€	78.87	15.00
133 VAL CG1	8.20	29.83	80.07	15.00
133 VAL CG2	6.37	30.65	78.66	15.00

134 ALA N	9.98	32.08	77.21	15.00
134 ALA CA	10.51	32.42	75.89	15.00
134 ALA C	10.71	31.08	75.20	15.00
134 ALA O	10.94	30.09	75.89	15.00
134 ALA CB	11.81	33.16	76.01	15.00
135 ILE N	10.58	31.02	73.88	15.00
135 ILE CA	10.74	29.75	73.17	15.00
135 ILE C	11.25	29.93	71.74	15.00
135 ILE O	11.54	31.04	71.29	15.00
135 ILE CB	9.39	29.00	73.05	15.00
135 ILE CG		29.93	72.47	15.00
135 ILE CG		28.39	74.37	15.00
135 ILE CD		29.26	72.13	15.00
136 ASP N	11.37	28.80	71.04	15.00
136 ASP CA	11.80	28.77	69.65	15.00
136 ASP C	10.54	28.58	68.80	15.00
136 ASP 0	10.02	27.47	68.67	15.00
136 ASP CB	12.79	27.62	69.36	15.00
136 ASP CG		27.57	67.89	15.00
136 ASP OD		28.61	67.20	15.00
136 ASP OD		26.49	67.41	15.00
137 ALA N	10.05	29.68	68.24	15.00
137 ALA CA	8.87	29.65	67.41	15.00
137 ALA C	9.23	29.70	65.91	15.00
137 ALA 0	8.38	30.00	65.07	15.00
137 ALA CE	7.97	30.81	67.79	15.00
138 SER N	10.47	29.35	65.57	15.00
138 SER CA	10.91	29.38	64.18	15.00
138 SER C	10.40	28.27	63.28	15.00
138 SER 0	10.47	28.39	62.06	15.00
138 SER CE	12.43	29.40	64.10	15.00
138 SER 00	12.93	30.59	64.68	15.00
139 LEU N	9.85	27.22	63.87	15.00
139 LEU C	9.36	26.07	63.12	15.00
139 LEU C	8.06	26.32	62.36	15.00
139 LEU O	7.07	26.74	62.94	15.00
139 LEU C	9.17	24.86	64.04	15.00
139 LEU C	g 10.27	24.49	65.02	15.00
139 LEU C	11.61	24.35	64.30	15.00
139 LEU C	10.33	25.54	66.14	15.00
140 THR N	8.06	26.00	61.06	15.00
140 THR C		26.19	60.24	15.00
140 THR C	5.73	25.38	60.86	15.00
140 THR 0	4.56	25.75	60.77	15.00

140	THR	CB	7.09	25.72	58.78	15.00
140	THR	OG1	8.35	26.21	58.30	15.00
140	THR	CG2	6.00	26.29	57.88	15.00
141	SER	N	6.11	24.29	61.54	15.00
141	SER	CA	5.14	23.42	62.20	15.00
141	SER	С	4.49	24.13	63.38	15.00
141	SER	0	3.32	23.91	63.69	15.00
141	SER	CB	5.83	22.12	62.64	15.00
141	SER		7.14	22.36	63.15	15.00
142	PHE		5.25	25.01	64.02	15.00
142	PHE		4.74	25.76	65.15	15.00
142	PHE	С	3.80	26.83	64.66	15.00
142	PHE	0	2.78	27.12	65.30	15.00
142	PHE	CB	5.88	26.41	65.94	15.00
142	PHE	CG	5.41	27.24	67.09	15.00
142	PHE		5.07	28.58	66.91	15.00
142	PHE	CD2	5.31	26.70	68.37	15.00
142	PHE	CE1	4.64	29.36	67.97	15.00
142	PHE	CE2	4.89	27.47	69.44	15.00
142	PHE	CZ	4.55	28.81	69.24	15.00
143	GLN	N	4.13	27.43	63.52	15.00
143	GLN	CA	3.31	28.49	62.99	15.00
143	GLN	С	1.93	28.03	62.57	15.00
143	GLN	0	0.95	28.72	62.84	15.00
143	GLN	CB	4.05	29.25	61.89	15.00
143	GLN	CG	5.12	30.14	62.48	15.00
143	GLN	CD	6.22	30.52	61.51	15.00
143	GLN		5.97	31.14	60.47	15.00
143	GLN	NE2	7.45	30.19	61.86	15.00
144	PHE		1.84	26.82	62.00	15.00
144	PHE		0.54	26.30	61.57	15.00
144	PHE	С	-0.14	25.30	62.52	15.00
144	PHE	0	-1.02	24.53	62.12	15.00
144	PHE	CB	0.56	25.79	60.11	15.00
144	PHE	CG	1.57	24.68	59.84	15.00
144	PHE	CD1	1.63	23.54	60.63	15.00
144	PHE	CD2	2.40	24.75	58.73	15.00
144	PHE	CE1	2.50	22.49	60.32	15.00
144	PHE	CE2	3.27	23.71	58.42	15.00
144	PHE	CZ	3.32	22.58	59.22	15.00
145	TYR	N	0.27	25.33	63.79	15.00
145	TYR	CA	-0.30	24.45	64.81	15.00
145	TYR	. C	-1.80	24.73	64.93	15.00
145		. 0	-2.22	25.89	64.92	1.5.00

		_		
145 TYR CB	0.36	24.72	66.17	15.00
145 TYR CG	-0.43	24.13	67.33	15.00
145 TYR CD1	-0.28	22.79	67.70	15.00
145 TYR CD2	-1.38	24.90	68.01	15.00
145 TYR CE1	-1.05	22.23	68.70	15.00
145 TYR CE2	-2.15	24.35	69.01	15.00
145 TYR CZ	-1.99	23.01	69.35	15.00
145 TYR OH	-2.76	22.46	70.34	15.00
146 SER N	-2.60	23.68	65.08	15.00
146 SER CA	-4.04	23.84	65.20	15.00
146 SER C	-4.65	22.98	66.30	15.00
146 SER 0	-5.72	23.29	66.81	15.00
146 SER CB	-4.74	23.59	63.84	15.00
146 SER OG	-4.46	22.30	63.31	15.00
147 LYS N	-3.96	21.91	66.67	15.00
147 LYS CA	-4.48	21.04	67.73	15.00
147 LYS C	-3.46	20.00	68.23	15.00
147 LYS 0	-2.52	19.67	67.51	15.00
147 LYS CB	-5.79	20.37	67.27	15.00
147 LYS CG	-5.76	19.70	65.88	15.00
147 LYS CD	-7.11	19.05	65.50	15.00
147 LYS CE	-7.19	18.65	64.01	15.00
147 LYS NZ	-7.28	19.82	63.07	15.00
148 GLY N	-3.63	19.55	69.48	15.00
148 GLY CA	-2.75	18.55	70.07	15.00
148 GLY C	-1.56	19.09	70.85	15.00
148 GLY 0	-1.43	20.29	71.06	15.00
149 VAL N	-0.71	18.18	71.31	15.00
149 VAL CA	0.50	18.54	72.07	15.00
149 VAL C	1.66	18.70	71.06	15.00
149 VAL O	2.06	17.74	70.39	15.00
149 VAL CB	0.83	17.46	73.14	15.00
149 VAL CG1	2.06	17.86	73.93	15.00
149 VAL CG2	-0.34	17.26	74.08	15.00
150 TYR N	2.21	19.90	70.99	15.00
150 TYR CA	3.25	20.21	70.03	15.00
150 TYR C	4.60	19.64	70.34	15.00
150 TYR O	5.26	20.08	71.28	15.00
150 TYR CB	3.39	21.72	69.86	15.00
150 TYR CG	4.42	22.11	68.81	15.00
150 TYR CD1	4.39	21.53	67.54	15.00
150 TYR CD2	5.44	23.02	69.11	15.00
150 TYR CE1	5.36	21.85	66.58	15.00
150 TYR CE2	6.41	23.33	68.15	15.00

150 TYR CZ	6.37	22.74	66.90	15.00
150 TYR OH	7.34	23.02	65.97	15.00
151 TYR N	5.03	18.69	69.53	15.00
151 TYR CA	6.35	18.11	69.70	15.00
151 TYR C	7.09	18.15	68.37	15.00
151 TYR O	6.65	17.56	67.39	15.00
151 TYR CB	6.30	16.68	70.20	15.00
151 TYR CG	7.67	16.22	70.63	15.00
151 TYR CD1	8.45	17.02	71.46	15.00
151 TYR CD2	8.20	15.02	70.17	15.00
151 TYR CE1	9.74	16.65	71.82	15.00
151 TYR CE2	9.50	14.63	70.52	15.00
151 TYR CZ	10.26	15.45	71.35	15.00
151 TYR OH	11.55	15.08	71.70	15.00
152 ASP N	8.21	18.85	68.35	15.00
152 ASP CA	8.98	18.96	67.12	15.00
152 ASP C	10.47	18.82	67.44	15.00
152 ASP 0	11.08	19.72	68.03	15.00
152 ASP CB	8.70	20.31	66.45	15.00
152 ASP CG	9.22	20.37	65.03	15.00
152 ASP OD1	8.48	19.96	64.12	15.00
152 ASP OD2	10.37	20.82	64.82	15.00
153 GLU N	11.03	17.68	67.05	15.00
153 GLU CA	12.44	17.38	67.31	15.00
153 GLU C	13.40	18.37	66.69	15.00
153 GLU O	14.59	18.34	66.98	15.00
153 GLU CB	12.76	15.94	66.86	15.00
153 GLU CG	12.29	15.56	65.44	15.00
153 GLU CD	13.28	15.95	64.33	15.00
153 GLU OE1	14.38	15.36	64.28	15.00
153 GLU OE2	12.95	16.84	63.50	15.00
154 SER N	12.89	19.26	65.85	15.00
154 SER CA	13.75	20.24	65.20	15.00
154 SER C	13.86	21.54	66.01	15.00
154 SER O	14.71	22.38	65.71	15.00
154 SER CB	13.20	20.54	63.80	15.00
154 SER OG	14.08	21.32	63.02	15.00
155 CYS N	13.02	21.69	67.03	15.00
155 CYS CA	13.01	22.90	67.86	15.00
155 CYS C	14.34	23.17	68.55	15.00 15.00
155 CYS 0	14.74	22.42	69.45	15.00
155 CYS CB	11.86	22.85	68.87	
155 CYS SG	11.05	24.47	69.10	15.00
156 ASN N	15.02	24.23	68.14	15.00

156	ASN	CA	16.33	24.57	68.70	15.00
156	ASN	С	16.29	25.23	70.07	15.00
156	ASN	0	16.21	26.46	70.19	15.00
156	ASN	CB	17.16	25.40	67.73	15.00
156	ASN	CG	18.54	25.71	68.28	15.00
156	ASN	OD1	18.89	26.86	68.48	15.00
156	ASN	ND2	19.32	24.66	68.57	15.00
157	SER	N	16.46	24.39	71.09	15.00
157	SER	CA	16.46	24.79	72.50	15.00
157	SER	С	17.39	25.95	72.87	15.00
157	SER	0	17.24	26.54	73.94	15.00
157	SER	CB	16.79	23.56	73.34	15.00
157	SER	OG	16.11	22.42	72.84	15.00
158	ASP	N	18.34	26.27	71.99	15.00
158	ASP	CA	19.29	27.35	72.24	15.00
158	ASP	С	18.88	28.70	71.64	15.00
158	ASP	0	19.31	29.76	72.09	15.00
158	ASP	CB	20.69	26.96	71.75	15.00
158	ASP	CG	21.57	26.39	72.87	15.00
158	ASP	OD1	21.02	25.82	73.85	15.00
158	ASP	OD2	22.82	26.53	72.78	15.00
159			18.07	28.66	70.59	15.00
159	ASN		17.63	29.89	69.97	15.00
159	ASN		16.22	30.22	70.46	15.00
159	ASN	0	15.25	29.58	70.04	15.00
159	ASN		17.69	29.76	68.45	15.00
159	ASN		16.80	30.75	67.73	15.00
159		OD1	16.93	31.98	67.88	15.00
159		ND2	15.86	30.22	66.96	15.00
160	LEU		16.13	31.16	71.39	15.00
160	LEU		14.83	31.58	71.93	15.00 15.00
160	LEU		14.48	32.83	71.16	15.00
160			15.25	33.79	71.14	15.00
160		CB	14.90	31.88	73.44	
	LEU		15.29	30.75	74.43	15.00 15.00
		CD1	15.63	31.34	75.78	15.00
		CD2	14.20	29.70	74.54	
	ASN		13.32	32.82	70.51	15.00
	ASN		12.91	33.94	69.69	15.00
	ASN		11.44	34.33	69.76	15.00
	ASN		10.94	34.97	68.85	15.00
	. ASN		13.25	33.62	68.24	15.00
	. ASN		12.59	32.35	67.78	15.00
161	. ASN	1 001	11.36	32.28	67.67	15.00

#### PCT/US96/17512 TABLE I 13.38 31.31 67.58 15.00 161 ASN ND2 33.90 70.78 15.00 10.72 162 HIS N 70.87 15.00 162 HIS CA 9.33 34.31 34.19 72.29 15.00 162 HIS C 8.84 33.09 72.84 15.00 162 HIS 0 8.74 69.91 15.00 162 HIS CB 8.45 33.51 34.12 69.68 15.00 162 HIS CG 7.10 35.47 69.42 15.00 162 HIS ND1 6.93 33.58 69.66 15.00 5.86 162 HIS CD2

168 GLY	CA	-0.99	26.12	83.13	15.00
168 GLY	С	-0.16	24.99	82.57	15.00
168 GLY	0	0.86	25.22	81.93	15.00
169 TYR	N	-0.61	23.78	82.81	15.00
169 TYR	CA	0.05	22.58	82.33	15.00
169 TYR	С	-1.02	21.51	82.42	15.00
169 TYR	0	-2.11	21.76	82.93	15.00
169 TYR	СВ	1.27	22.22	83.20	15.00
169 TYR	CG	1.02	22.11	84.70	15.00
169 TYR	CD1	0.52	20.94	85.28	15.00
169 TYR	CD2	1.32	23.18	85.56	15.00
169 TYR	CE1	0.32	20.84	86.66	15.00
169 TYR	CE2	1.12	23.08	86.94	15.00
169 TYR	CZ	0.62	21.91	87.48	15.00
169 TYR	OH	0.44	21.83	88.85	15.00
170 GLY	N	-0.71	20.33	81.90	15.00
170 GLY	CA	-1.65	19.23	81.94	15.00
170 GLY	С	-1.16	18.11	81.05	15.00
170 GLY	0	-0.03	18.14	80.54	15.00
171 ILE	N	-2.04	17.15	80.81	15.00
171 ILE	CA	-1.78	15.98	79.97	15.00
171 ILE	С	-2.96	15.87	79.00	15.00
171 ILE	0	-3.95	16.60	79.14	15.00
171 ILE	CB	-1.62	14.73	80.86	15.00
171 ILE	CG1	-1.47	13.46	80.02	15.00
171 ILE	CG2	-2.76	14.65	81.87	15.00
171 ILE	CD1	-1.03	12.22	80.81	15.00
172 GLN	N	-2.86	15.03	77.96	15.00
172 GLN	CA	-3.97	14.90	77.01	15.00
172 GLN	C	-4.16	13.50	76.45	15.00
172 GLN	0	-5.04	12.76	76.88	15.00
172 GLN	CB	-3.79	15.89	75.87	15.00
172 GLN	CG	-4.95	15.94	74.92	15.00
172 GLN	CD	-4.83	17.08	73.95	15.00
172 GLN	OE1	-3.81	17.24	73.28	15.00
172 GLN	NE2	-5.86	17.91	73.88	15.00
173 LYS	N	-3.39	13.15	75.44	15.00
173 LYS	CA	-3.51	11.81	74.89	15.00
173 LYS	C	-2.36	11.09	75.55	15.00
173 LYS	0	-1.55	10.43	74.90	15.00
173 LYS	CB	-3.36	11.82	73.36	15.00
173 LYS	CG	-4.57	12.39	72.60	15.00
173 LYS	CD	-5.81	11.53	72.82	15.00
173 LYS	CE	-7.06	12.17	72.19	15.00

	٠	TABLE I		
173 LYS NZ	-8.35	11.45	72.50	15.00
174 GLY N	-2.29	11.24	76.87	15.00
174 GLY CA	-1.20	10.64	77.60	15.00
174 GLY C	0.03	11.49	77.36	15.00
174 GLY 0	1.17	11.08	77.62	15.00
175 ASN N	-0.19	12.72	76.92	15.00
175 ASN CA	0.93	13.59	76.64	15.00
175 ASN C	1.02	14.78	77.57	15.00
175 ASN 0	0.07	15.56	77.70	15.00
175 ASN CB	0.88	14.04	75.18	15.00
175 ASN CG	0.72	12.87	74.20	15.00
175 ASN OD1	-0.28	12.77	73.49	15.00
175 ASN ND2	1.71	11.97	74.19	15.00
176 LYS N	2.15	14.88	78.27	15.00
176 LYS CA	2.42	15.97	79.20	15.00
176 LYS C	2.40	17.22	78.34	15.00
176 LYS 0	2.76	17.17	77.16	15.00
176 LYS CB	3.83	15.84	79.78	15.00
176 LYS CG	4.38	14.45	79.93	15.00
176 LYS CD	4.27	13.97	81.36	15.00
176 LYS CE	4.98	12.63	81.55	15.00
176 LYS NZ	4.97	12.14	82.96	15.00
177 HIS N	2.01	18.36	78.91	15.00
177 HIS CA	1.99	19.58	78.13	15.00
177 HIS C	1.88	20.86	78.95	15.00
177 HIS O	1.47	20.84	80.12	15.00
177 HIS CB	0.88	19.53	77.06	15.00 15.00
177 HIS CG	-0.52	19.66	77.59	15.00
177 HIS ND1		18.63	77.56 78.16	15.00
177 HIS CD2	-1.16	20.71	78.18	15.00
177 HIS CE1		19.03 20.29	78.45	15.00
177 HIS NE2		21.96	78.43	15.00
178 TRP N	2.25	23.29	78.89	15.00
178 TRP CA	2.17	24.04		15.00
178 TRP C	1.16	23.95		15.00
178 TRP 0	1.20 3.52	23.95	78.78	15.00
178 TRP CB	4.66	23.32	79.46	15.00
178 TRP CG	4.00	23.34	79.40	15.00

22.70

23.34 22.35

22.73

23.83

22.60

5.71

4.94

6.65

6.21

4.26

6.80

178 TRP CD1

178 TRP CD2

178 TRP NE1

178 TRP CE2

178 TRP CE3

178 TRP CZ2

78.87

80.86

79.80

81.03

81.99

82.29

15.00

15.00

15.00

15.00

15.00

15.00

178	TRP	CZ3	4.85	23.70	83.23	15.00
178	TRP	CH2	6.12	23.09	83.38	15.00
179	ILE	N	0.22	24.75	78.66	15.00
179	ILE	CA	-0.79	25.51	77.92	15.00
179	ILE	С	-0.20	26.89	77.64	15.00
179	ILE	0	-0.07	27.71	78.54	15.00
179	ILE	СВ	-2.08	25.66	78.74	15.00
179	ILE	CG1	-2.67	24.27	79.03	15.00
179	ILE	CG2	-3.07	26.56	78.02	15.00
179	ILE	CD1	-3.80	24.25	80.04	15.00
180	ILE	N	0.21	27.09	76.39	15.00
180	ILE	CA	0.83	28.33	75.96	15.00
180	ILE	С	-0.17	29.28	75.34	15.00
180	ILE	0	-1.09	28.85	74.65	15.00
180	ILE	CB	1.99	28.05	74.99	15.00
180	ILE	CG1	3.32	28.14	75.72	15.00
180	ILE	CG2	1.97	28.97	73.83	15.00
180	ILE	CD1	3.53	27.06	76.73	15.00
181	LYS	N	0.00	30.57	75.64	15.00
181	LYS	CA	-0.85	31.66	75.14	15.00
181	LYS	С	-0.01	32.55	74.22	15.00
181	LYS	0	0.91	33.23	74.68	15.00
181	LYS	CB	-1.38	32.51	76.30	15.00
181	LYS	CG	-2.27	33.66	75.84	15.00
181	LYS		-2.51	34.72	76.91	15.00
181	LYS		-3.38	35.84	76.37	15.00
181	LYS		-3.58	36.94	77.35	15.00
182			-0.32	32.56	72.93	15.00
182	ASN		0.44	33.36	71.97	15.00
182			-0.18	34.74	71.73	15.00
182			-1.29	35.00	72.17	15.00
182	asn		0.55	32.59	70.66	15.00
182			1.83	32.87	69.93	15.00
		OD1	2.41	33.95	70.05	15.00
		ND2	2.31	31.89	69.17	15.00 15.00
183			0.55	35.64	71.08	
183		. CA	0.03	36.98	70.82	15.00 15.00
183			-0.32	37.19	69.34	
183			-0.25	33.31	68.82	15.00
183		CB	1.03	38.04	71.25	15.00
183		OG	2.31	37.20	70.69	15.00
184			-0.74	36.12	68.68	15.00
184		CA	-1.05	36.18	67.26	15.00
184	TRE	C	-2.53	36.23	66.88	15.00

184	TRP	0	-2.87	36.09	65.71	15.00
184	TRP	CB	-0.36	35.01	66.55	15.00
184	TRP	CG	1.12	35.12	66.54	15.00
184	TRP	CD1	1.85	36.23	66.82	15.00
184	TRP	CD2	2.05	34.09	66.23	15.00
184	TRP	NE1	3.18	35.97	66.69	15.00
184	TRP	CE2	3.34	34.66	66.33	15.00
184	TRP	CE3	1.94	32.75	65.87	15.00
184	TRP	CZ2	4.51	33.92	66.09	15.00
184	TRP	CZ3	3.10	32.01	65.63	15.00
184	TRP	CH2	4.37	32.60	65.74	15.00
185	GLY	N	-3.40	36.48	67.85	15.00
185	GLY	CA	-4.82	36.54	67.56	15.00
185	GLY	С	-5.46	35.18	67.75	15.00
185	GLY	0	-4.75	34.16	67.80	15.00
186	GLU	N	-6.78	35.15	67.86	15.00
186	GLU	CA	-7.54	33.91	68.07	15.00
186	GLU	C	-7.53	32.99	66.85	15.00
186	GLU	0	-7.65	31.78	67.00	15.00
	GLU		-8.98	34.24	68.44	15.00
	GLU		-9.10	35.32	69.49	15.00
	GLU		-10.45	36.03	69.45	15.00
186	GLU	OE1	-10.66	36.84	68.51	15.00
186	GLU	OE2	-11.29	35.77	70.34	15.00
187	ASN	N	-7.43	33.58	65.66	15.00
	ASN		-7.43	32.78	64.42	15.00
	ASN		-6.14	31.97	64.30	15.00
	ASN		-6.07	31.05	63.49	15.00
	ASN		-7.64	33.68	63.19	15.00
	ASN		-8.60	33.05	62.14	15.00
	ASN		-8.17	32.52	61.10	15.00
	ASN		-9.90	33.18	62.38	15.00
	TRP		-5.12	32.31	65.10	15.00
	TRP		-3.87	31.57	65.07	15.00
188	TRP		-4.06	30.34	65.95	15.00
188			-4.78	30.41	66.95	15.00
	TRP		-2.71	32.38	65.64	15.00
188			-1.45	31.59	65.56	15.00
188		CD1	-0.68	31.41	64.46	15.00
188		CD2	-0.89	30.74	66.58	15.00
188	TRP		0.30	30.49	64.71	15.00
188	TRP		0.20	30.06	66.00	15.00
188		CE3	-1.21	30.49	67.92	15.00
138	TRP	CZ2	0.97	29.14	66.71	15.00

188	TRP	CZ3	-0.44	29.57	68.62	15.00
188	TRP	CH2	0.64	28.92	68.02	15.00
189	GLY	N	-3.38	29.25	65.60	15.00
189	GLY	CA	-3.46	28.02	66.36	15.00
189	GLY	c ·	-4.84	27.67	66.87	15.00
189	GLY	0	-5.86	27.96	66.24	15.00
190	ASN	N	-4.87	27.07	68.04	15.00
190	ASN	CA	-6.10	26.65	68.68	15.00
190	ASN	С	-6.74	27.83	69.39	15.00
190	ASN	0	-6.67	27.92	70.61	15.00
190	ASN	CB	-5.76	25.58	69.71	15.00
190	ASN	CG	-6.97	24.84	70.19	15.00
190	ASN	OD1	-8.11	25.26	69.96	15.00
190	ASN	ND2	-6.73	23.72	70.87	15.00
191	LYS	N	-7.33	28.75	68.61	15.00
191	LYS	CA	-7.97	29.96	69.14	15.00
191	LYS	С	-7.01	30.82	69.95	15.00
191	LYS	0	-7.36	31.36	71.00	15.00
191	LYS	CB	-9.20	29.62	69.99	15.00
191	LYS	CG	-10.33	28.92	69.26	15.00
191	LYS	CD	-11.50	28.66	70.20	15.00
191	LYS	CE	-12.26	27.39	69.82	15.00
191	LYS	NZ	-11.42	26.16	70.01	15.00
192	GLY	N	-5.79	30.96	69.45	15.00
192	GLY	CA	-4.80	31.75	70.13	15.00
192	GLY	С	-3.88	30.93	70.99	15.00
192	GLY	0	-2.79	31.39	71.32	15.00
193	TYR	N	-4.28	29.72	71.35	15.00
193	TYR		-3.43	28.86	72.18	15.00
193	TYR		-2.80	27.69	71.45	15.00
193	TYR	0	-3.25	27.27	70.39	15.00
193	TYR		-4.21	28.31	73.39	15.00
193	TYR		-4.63	29.36	74.36	15.00
193		CD1	-3.77	29.78	75.37	15.00
	TYR		-5.85	30.00	74.22	15.00
193		CE1	-4.12	30.82	76.22	15.00
193	TYR	CE2	-6.21	31.05	75.06	15.00
193			-5.34	31.45	76.06	15.00
193	TYR		-5.68	32.50	76.87	15.00
194	ILE		-1.73	27.17	72.05	15.00
194	ILE		-1.01	26.02	71.53	15.00
194	ILE		-0.53	25.21	72.71	15.00
194	ILE	0	0.04	25.75	73.66	15.00
194	ILE	CB	0.20	26.40	70.62	15.00

194	ILE	CG1	1.05	25.15	70.32	15.00
194	ILE	CG2	1.04	27.49	71.24	15.00
194	ILE	CD1	2.33	25.42	69.60	15.00
195	LEU	N	-0.86	23.92	72.69	15.00
195	LEU	CA	-0.45	23.00	73.73	15.00
195	LEU	С	0.90	22.46	73.34	15.00
195	LEU	0	0.99	21.68	72.42	15.00
195	LEU	CB	-1.43	21.83	73.83	15.00
195	LEU	CG	-2.45	21.90	74.96	15.00
195	LEU	CD1	-3.38	23.07	74.76	15.00
195	LEU	CD2	-3.23	20.60	74.98	15.00
196	MET	N	1.95	22.94	74.01	15.00
196	MET	CA	3.31	22.50	73.75	15.00
196	MET	C	3.70	21.30	74.64	15.00
196	MET	0	2.97	20.96	75.57	15.00
196	MET	CB	4.27	23.66	73.96	15.00
196	MET	CG	4.16	24.69	72.88	15.00
196	MET	SD	5.11	26.17	73.23	15.00
	MET		6.79	25.64	72.90	15.00
	ALA		4.85	20.69	74.38	15.00
	ALA		5.27	19.52	75.15	15.00
	ALA		5.99	19.75	76.49	15.00
	ALA		7.03	20.42	76.57	15.00
	ALA		6.07	18.58	74.28	15.00
	ARG		5.43	19.14	77.54	15.00
	ARG		5.99	19.22	78.88	15.00
	ARG		6.80	17.96	79.20	15.00
198	ARG		6.41	16.84	78.86	15.00
198	ARG		4.89	19.44	79.92	15.00
	ARG		5.35	19.32	81.39	15.00
	ARG		4.48	20.12	82.35	15.00
	ARG		3.08	19.72	82.36	15.00
	ARG		2.62	18.64	82.99	15.00
198		NH1	3.46	17.86	83.66	15.00
	ARG		1.33	18.34	82.95	15.00
199			7.96	18.18	79.83	15.00
	ASN		8.87	17.11	80.24	15.00
	asn		9.57	16.30	79.13	15.00
	ASN		10.36	15.40	79.43	15.00
199			8.21	16.18	81.28	15.00
199			8.15	16.81	82.68	15.00
199		OD1	7.06	17.06	83.21	15.00
199		ND2	9.31	17.07	83.27	15.00
200	LYS	N	9.31	16.63	77.88	15.00

200	TVC	Ca	0 07	15 04	26 22	15 00
200	LYS LYS		9.97 11.29	15.94 16.67	76.77 76.54	15.00 15.00
200	LYS		11.54	17.17	75.45	15.00
200	LYS		9.11	15.96	75.49	15.00
200	LYS		8.09	14.81	75.36	15.00
200	LYS		8.32	14.01	74.06	15.00
200	LYS		7.22	12.96	73.78	15.00
200	LYS		5.87	13.49	73.78	15.00
201			12.13	16.71	77.57	15.00
	ASN		13.42	17.39	77.56	15.00
201			13.27	18.81	77.00	15.00
	ASN		13.24	18.99	75.80	15.00
	ASN		14.47	16.61	76.74	15.00
	ASN		15.92	17.21	76.86	15.00
	ASN		16.88	16.46	77.06	15.00
	ASN		16.06	18.52	76.67	15.00
202			13.15	19.80	77.89	15.00
	ASN		13.04	21.22	77.54	15.00
	ASN		12.73	21.51	76.05	15.00
	ASN		13.56	22.10	75.34	15.00
	ASN		14.35	21.92	77.95	15.00
	ASN		14.13	23.30	78.56	15.00
	ASN		13.04	23.64	79.00	15.00
	ASN		15.18	24.09	78.60	15.00
	ALA		11.55	21.12	75.59	15.00
	ALA		11.67	20.72	76.05	15.00
203	ALA		11.15	21.27	74.19	15.00
203	ALA	•	10.97	20.66	73.61	15.00
203	ALA		11.04	22.76	73.85	15.00
203	ALA		10.23	23.49	74.41	15.00
	CYS		11.83	23.20	72.87	15.00
	CYS		11.81	24.59	72.40	15.00
204	CYS		12.38	25.57	73.42	15.00
•	CYS		12.06	26.76	73.36	15.00
	CYS		10.39	25.02	72.00	15.00
204		SG	9.66	24.16	70.56	15.00
205	GLY		13.21	25.07	74.33	15.00
205	GLY		13.84	25.89	75.36	15.00
205	GLY		12.87	26.70	76.20	15.00
205	GLY		13.17	27.80	76.64	15.00
206	ILE		11.72	26.10	76.48	15.00
206	ILE		10.67	26.74	77.23	15.00
206	ILE		11.05	27.11	78.67	15.00
206	ILE		10.90	28.27	79.08	15.00
200		9	_0.50			

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				TABLE I		•
206	ILE	СВ	9.41	25.86	77.17	15.00
206	ILE	CG1	8.20	26.58	77.77	15.00
206	ILE	CG2	9.69	24.53	77.81	15.00
206	ILE	CD1	6.88	25.88	77.49	15.00
207	ALA	N	11.62	26.17	79.41	15.00
207	ALA	CA	12.03	26.40	80.79	15.00
207	ALA	C	13.38	27.11	80.93	15.00
207	ALA	0	13.98	27.10	82.01	15.00
207	ALA	CB	12.05	25.09	81.54	15.00
208	ASN	N	13.84	27.74	79.86	15.00
208	ASN	CA	15.12	28.43	79.86	15.00
208	ASN	С	15.08	29.92	80.08	15.00
208	ASN	0	16.13	30.53	80.28	15.00
208	ASN	CB	15.84	28.22	78.54	15.00
208	ASN	CG	16.81	27.09	78.60	15.00
208	ASN	OD1	16.50	26.04	79.14	15.00
208	ASN	ND2	17.99	27.30	78.04	15.00
209	LEU	N	13.91	30.54	79.98	15.00
209	LEU	CA	13.84	31.98	80.18	15.00
209	LEU	C	12.65	32.45	81.02	15.00
209	LEU	Ö	12.23	33.63	80.94	15.00
209	LEU	CB	13.89	32.69	78.83	15.00
	LEU		14.33	34.16	78.81	15.00
	LEU		15.60	34.32	79.63	15.00
	LEU		14.56	34.61	77.37	15.00
	ALA		12.15	31.57	81.87	15.00
	ALA		12.50	30.94	81.70	15.00
	ALA		10.99	31.87	82.71	15.00
	ALA		10.41	31.04	83.16	15.00
	ALA		11.33	33.00	83.69	15.00
	ALA		12.46	33.21	84.13	15.00
	SER		10.26	33.74	84.06	15.00
211			10.37	34.83	85.02	15.00
211			8.96	35.33	85.31	15.00
	SER		8.09	35.24	84.45	15.00 15.00
	SER		11.22	35.98	84.46	15.00
211			10.50	36.76	83.53	
212			8.72	35.78	86.53	15.00
212			7.42	36.30	86.88 87.78	15.00 15.00
	PHE		7.64	37.49		15.00
212			8.68	37.61	88.42	15.00
212			6.57	35.23	87.56 88.75	15.00
212			7.23	34.60	89.94	15.00
212	PHE	CD1	7.39	35.31	07.74	10.00

212	PHE	CD2	7.68	33.28	88.69	15.00
212	PHE	CE1	7.99	34.73	91.05	15.00
212	PHE	CE2	8.28	32.68	89.78	15.00
212	PHE	CZ	8.44	33.41	90.97	15.00
213	PRO	N	6.69	38.43	87.80	15.00
213	PRO	CA	6.84	39.62	88.65	15.00
213	PRO	С	6.38	39.34	90.09	15.00
213	PRO	0	5.56	38.44	90.33	15.00
213	PRO	CB	5.93	40.62	87.97	15.00
213	PRO	CG	4.79	39.75	87.52	15.00
213	PRO	CD	5.47	38.52	86.97	15.00
214	LYS	N	6.96	40.05	91.04	15.00
214	LYS	CA	6.57	39.89	92.42	15.00
214	LYS	С	5.65	41.06	92.68	15.00
214	LYS	0	6.12	42.17	92.91	15.00
214	LYS	CB	7.78	39.94	93.36	15.00
214	LYS	CG	8.68	38.72	93.27	15.00
214	LYS	CD	9.78	38.70	94.34	15.00
214	LYS	CE	10.67	39.94	94.26	15.00
214	LYS	NZ	11.94	39.80	95.03	15.00
215	MET	N	4.36	40.85	92.51	15.00
215	MET	CA	3.39	41.91	92.77	15.00
215	MET	С	2.86	41.71	94.19	15.00
215	MET	CB	2.25	41.86	91.76	15.00
215	MET	CG	1.06	42.74	92.12	15.00
215	MET	SD	-0.32	42.54	90.98	15.00
215	MET	CE	0.28	43.53	89.53	15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (Å<sup>2</sup>) for the cathepsin K complex with inhibitor 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone.

Residue	Atom	X	Y	Z	В
1 ALA	СВ	~54.11	-32.66	67.34	15.00
1 ALA	C	-54.02	-32.71	64.82	15.00
1 ALA	0	-53.62	-33.85	64.58	15.00
1 ALA	N	-56.02	-33.44	65.99	15.00
1 ALA	CA	-54.90	-32.46	66.05	15.00
2 PRO	N	-53.80	-31.67	63.99	15.00
2 PRO	CD	-54.47	-30.37	64.11	15.00
2 PRO	CA	-52.98	-31.72	62.76	15.00
2 PRO	CB	-53.14	-30.31	62.20	15.00
2 PRO	CG	-54.52	-29.90	62.67	15.00
2 PRO		-51.51	-32.06	62.95	15.00
2 PRO	0	-50.99	-32.06	64.08	15.00
3 ASP		-50.81	-32.33	61.85	15.00
3 ASP		-49.39	-32.67	61.91	15.00
3 ASP		-49.04	-33.71	60.85	15.00
3 ASP	CG	-49.27	-35.14	61.32	15.00
3 ASP	OD1	-48.47	-35.61	62.17	15.00
3 ASP		-50.24	-35.79	60.85	15.00
3 ASP	С	-48.52	-31.44	61.73	15.00
3 ASP	0	-47.93	-31.24	60.68	15.00
4 SER	N	-48.41	-30.64	62.79	15.00
4 SER	CA	-47.62	-29.41	62.75	15.00
4 SER	CB	-48.55	-28.25	62.37	15.00
4 SER	OG	-47.82	-27.08	62.01	15.00
4 SER	C	-47.01	-29.13	64.12	15.00
4 SER		-47.59	-29.49	65.15	15.00
5 VAL	N	-45.83	-28.51	64.15	15.00
5 VAL	CA	-45.19	-28.15	65.41	15.00
5 VAL	CB	-44.36	-29.31	66.03	15.00
5 VAL		-43.06	-29.53	65.29	15.00
5 VAL	CG2	-44.08	-29.04	67.50	15.00
5 VAL	С	-44.34	-26.88	65.24	15.00
5 VAL	0	-43.64	-26.69	64.25	15.00
6 ASP	N .	-44.48	-25.96	66.20	15.00

6	ASP	CA	-43.78	-24.69	66.19	15.00
6	ASP	CB	-44.75	-23.59	65.74	15.00
6	ASP	CG	-44.11	-22.19	65.67	15.00
6	ASP	OD1	-42.99	-21.99	66.18	15.00
6	ASP	OD2	-44.75	-21.27	65.13	15.00
6	ASP	С	-43.37	-24.47	67.64	15.00
6	ASP	0	-44.19	-24.09	68.48	15.00
7	TYR	N	-42.10	-24.68	67.95	15.00
7	TYR	CA	-41.65	-24.50	69.33	15.00
7	TYR	CB	-40.30	-25.18	69.53	15.00
7	TYR	CG	-40.41	-26.69	69.53	15.00
7	TYR	CD1	-40.91	-27.37	70.64	15.00
7	TYR	CE1	-41.02	-28.74	70.65	15.00
7	TYR	CD2	-40.02	-27.43	68.42	15.00
7	TYR	CE2	-40.13	-28.80	68.42	15.00
7	TYR	CZ	-40.63	-29.45	69.53	15.00
7	TYR	OH	-40.70	-30.82	69.53	15.00
7	TYR	С	-41.62	-23.07	69.82	15.00
7	TYR	0	-41.41	-22.81	71.00	15.00
8	ARG	N	-41.83	-22.12	68.92	15.00
8	ARG	CA	-41.84	-20.72	69.31	15.00
8	ARG	CB	-42.00	-19.80	68.09	15.00
8	ARG	CG	-40.82	-19.80	67.14	15.00
8	ARG	CD	-41.13	-18.98	65.91	15.00
8	ARG	NE	-42.05	-19.66	65.00	15.00
8	ARG	CZ	-42.68	-19.07	64.00	15.00
8	ARG	NH1	-42.49	-17.78	63.77	15.00
8	ARG	NH2	-43.50	-19.77	63.22	15.00
8	ARG	С	-43.00	-20.51	70.28	15.00
8	ARG	0	-42.87	-19.79	71.28	15.00
9	LYS	N	-44.10	-21.19	70.00	15.00
9	LYS	CA	-45.30	-21.10	70.82	15.00
9	LYS	CB	-46.49	-21.67	70.05	15.00
9	LYS	CG	-46.76	-21.07	68.69	15.00
9	LYS	CD	-48.04	-21.67	68.14	15.00
9	LYS	CE	-48.28	-21.36	66.69	15.00
9	LYS	NZ	-49.49	-22.07	66.19	15.00
9	LYS	С	-45.20	-21.83	72.16	15.00
9	LYS	0	-46.13	-21.78	72.97	15.00
0	LYS	N	-44.10	-22.53	72.40	15.00
0	LYS	CA	-43.92	-23.27	73.64	15.00
0	LYS	CB	-43.47	-24.71	73.36	15.00
0	LYS	CG	-43.75	-25.23	71.96	15.00

10	LYS	CD	-45.22	-25.49	71.73	15.00
10	LYS (	CE	-45.52	-26.97	71.83	15.00
10	LYS I	NZ	-45.21	-27.50	73.19	15.00
10	LYS (	С	-42.88	-22.63	74.54	15.00
10	LYS (	0	-42.67	-23.06	75.67	15.00
11	GLY 1	N	-42.16	-21.63	74.03	15.00
11	GLY (	CA	-41.15	-20.98	74.83	15.00
11	GLY (	C	-39.79	-21.63	74.68	15.00
11	GLY (	0	-38.90	-21.42	75.50	15.00
12	TYR I	N	-39.60	-22.38	73.60	15.00
12	TYR (	CA	-38.34	-23.07	73.36	15.00
12	TYR (	CB	-38.56	-24.39	72.61	15.00
12	TYR (	CG	-39.12	-25.53	73.43	15.00
12	TYR (	CD1	-40.27	-25.37	74.19	15.00
12	TYR (	CE1	-40.82	-26.43	74.89	15.00
12	TYR	CD2	-38.53	-26.79	73.39	15.00
12	TYR	CE2	-39.08	-27.86	74.09	15.00
12	TYR	CZ	-40.23	-27.67	74.84	15.00
12	TYR (	OH	-40.79	-28.71	75.55	15.00
12	TYR (	С	-37.31	-22.26	72.60	15.00
12	TYR	0	-36.14	-22.61	72.57	15.00
13	VAL 1	N	-37.70	-21.18	71.93	15.00
13	VAL	CA	-36.72	-20.42	71.18	15.00
13	VAL	CB	-36.97	-20.48	69.67	15.00
13	VAL	CG1	-37.21	-21.91	69.23	15.00
13	VAL	CG2	-38.14	-19.62	69.30	15.00
13	VAL	С	-36.63	-18.98	71.64	15.00
13	VAL	0	-37.62	-18.38	72.02	15.00
14	THR	N	-35.41	-18.45	71.65	15.00
14	THR	CA	-35.16	-17.08	72.07	15.00
	THR		-33.75	-16.96	72.66	15.00
	THR		-32.79	-17.40	71.71	15.00
14	THR	CG2	-33.63	-17.80	73.91	15.00
14	THR	С	-35.32	-16.08	70.91	15.00
14	THR		-35.57	-16.48	69.77	15.00
15			-35.24	-14.77	71.20	15.00
15			-35.20	-14.14	72.53	15.00
15			-35.37	-13.75	70.15	15.00
15	PRO	CB	-35.25	-12.44	70.93	15.00
15	PRO	CG	-35.83	-1.2.79	72.26	15.00
15	PRO	С	-34.26	-13.87	69.11	15.00
15	PRO	0	-33.13	-14.23	69.44	15.00
16	VAL	N	-34.59	-13.55	67.85	15.00

16	VAL	CA	-33.64	-13.61	66.75	15.00
16	VAL	CB	-34.33	-13.31	65.40	15.00
16	VAL	CG1	-33.34	-13.41	64.25	15.00
16	VAL	CG2	-35.47	-14.26	65.19	15.00
16	VAL	C	-32.48	-12.65	66.96	15.00
16	VAL	0	-32.69	-11.49	67.32	15.00
17	LYS	N	-31.28	-13.13	66.71	15.00
17	LYS	CA	-30.07	-12.35	66.86	15.00
17	LYS	CB	-29.12	-13.04	67.84	15.00
17	LYS	CG	-29.68	-13.07	69.24	15.00
17	LYS	CD	-28.91	-14.03	70.13	15.00
17	LYS	CE	-29.54	-14.07	71.53	15.00
17	LYS	NZ	-31.02	-14.22	71.43	15.00
17	LYS	C	-29.39	-12.14	65.52	15.00
17	LYS	0	-29.72	-12.79	64.54	15.00
18	ASN	N	-28.42	-11.23	65.50	15.00
18	ASN	CA	-27.68	-10.87	64.29	15.00
18	ASN	CB	-27.77	-9.36	64.10	15.00
18	asn	CG	-27.26	-8.90	62.75	15.00
18	ASN	OD1	-26.16	-9.25	62.32	15.00
18	ASN	ND2	-28.07	-8.10	62.08	15.00
18	asn	С	-26.23	-11.28	64.45	15.00
18	asn	0	-25.55	-10.79	65.35	15.00
19	GLN	N	-25.72	-12.16	63.60	15.00
19	GLN	CA	-24.33	-12.56	63.73	15.00
19	GLN	CB	-24.00	-13.81	62.90	15.00
19	GLN	CG	-24.44	-13.80	61.45	15.00
19	GLN	CD	-24.06	-15.09	60.72	15.00
19	GLN	OE1	-24.91	-15.79	60.16	15.00
	GLM		-22.77	-15.41	60.71	15.00
19	GLN		-23.40	-11.40	63.39	15.00
19	GLN		-22.27		63.88	15.00
	GLY		-23.91	-10.46	62.60	15.00
	GLY		-23.14	-9.30	62.21	15.00
	GLY			-9.67		
	GLY		-22.39			
	GLN		-20.93		61.2 <b>6</b>	15.00
	GLN		-19.85	-9.33	60.32	15.00
	GLN		-19.08	-8.03		15.00
		CG	-19.94		59.8 <b>4</b>	
		CD	-20.87			
			-20.43		57.49	
21	GLN	NE2	-22.17	-6.95	58.90	15.00

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21 GLN C	-18.91	-10.35	60.98	15.00
21 GLN O	-17.86	-10.01	61.52	15.00
22 CYS N	-19.31	-11.61	60.97	15.00
22 CYS CA	-18.53	-12.68	61.58	15.00
22 CYS C	-19.27	-13.95	61.19	15.00
22 CYS O	-20.48	-14.03	61.36	15.00
22 CYS CB	-18.50	-12.50	63.11	15.00
22 CYS SG	-17.85	-13.86	64.14	15.00
23 GLY N	-18.56	-14.88	60.56	15.00
23 GLY CA	-19.19	-16.12	60.13	15.00
23 GLY C	-19.43	-17.05	61.30	15.00
23 GLY O	-18.94	-18.17	61.31	15.00
24 SER N	-20.24	-16.60	62.25	15.00
24 SER CA	-20.56	-17.36	63.44	15.00
24 SER CB	-20.46	-16.45	64.66	15.00
24 SER OG	-21.23	-15.30	64.42	15.00
24 SER C	-21.95	-17.97	63.37	15.00
24 SER O	-22.54	-18.31	64.40	15.00
25 CYS N	-22.49	-18.14	62.17	15.00
25 CYS CA	-23.81	-18.74	62.02	15.00
25 CYS CB	-24.21	-18.82	60.54	15.00
25 CYS SG	-23.17	-19.85	59.47	15.00
25 CYS C	-23.88	-20.12	62.68	15.00
25 CYS O	-24.96	-20.59	63.04	15.00
25 INH C1	-26.94	-9.70	58.69	15.00
25 INH C2	-26.28	-10.48	59.65	15.00
25 INH C3	-25.12	-11.19	59.30	15.00
25 INH C4	-24.61	-11.12	58.00	15.00
25 INH C5	-25.28	-10.33	57.05	15.00
25 INH C6	-26.44	-9.62	57.39	15.00
25 INH C7	-23.37	-11.90	57.62	15.00
25 INH 08	-23.43	-13.32	57.82	15.00
25 INH C9	-22.85	-14.36	57.02	15.00
25 INH 010	-21.63	-14.58	56.99	15.00
25 INH C11	-23.27	-16.14	55.41	15.00
25 INH C12	-22.06	15.67	54.58	15.00
25 INH C13		-15.18	53.14	15.00
25 INH C14		-15.22	52.56	15.00
25 INH C15		-16.04	52.32	15.00
25 INH C1.6		-17.51	56.01	15.00
25 INH 017		-18.50	55.66	15.00
25 INH N18		-17.60	56.86	
25 INH C19	-21.48	-18.89	57.42	15.00

25 INH C20	-20.01	-19.11	57.02	15.00
25 INH C21	-19.59	-19.34	55.56	15.00
25 INH C22	-19.45	-20.84	55.30	15.00
25 INH C23	-18.25	-18.64	55.30	15.00
25 INH N24	-23.71	-15.07	56.30	15.00
25 INH C25	-21.62	-19.06	58.94	15.00
25 INH 026	-21.55	-17.95	59.50	15.00
25 INH C27	-20.53	-20.00	59.45	15.00
25 INH 028	-20.36	-21.26	58.72	15.00
25 INH C29	-19.70	-22.40	59.29	15.00
25 INH C30	-19.53	-23.60	58.35	15.00
25 INH C31	-20.80	-24.42	58.08	15.00
26 TRP N	-22.73	-20.75	62.90	15.00
26 TRP CA	-22.65	-22.06	63.54	15.00
26 TRP CB	-21.30	-22.75	63.25	15.00
26 TRP CG	-20.09	-22.04	63.79	15.00
26 TRP CD2	-19.48		65.08	15.00
26 TRP CE2	-18.36	-21.38	65.14	15.00
26 TRP CE3	-19.77	-23.04	66.19	15.00
26 TRP CD1	-19.33	-21.11	63.15	15.00
26 TRP NE1	-18.29	-20.70	63.95	15.00
26 TRP CZ2	-17.53	-21.31	66.27	15.00
26 TRP CZ3	-18.94	-22.97	67.31	15.00
26 TRP CH2	-17.83	-22.11	67.33	15.00
26 TRP C	-22.89	-22.02	65.06	15.00
26 TRP O	-23.59	-22.87	65.61	15.00
27 ALA N	-22.34	-21.01	65.75	15.00
27 ALA CA	-22.49	-20.86	67.19	15.00
27 ALA CB	-21.58	-19.78	67.71	15.00
27 ALA C	-23.93	-20.53	67.54	15.00
27 ALA O	-24.46	-20.98	68.55	15.00
28 PHE N	-24.55	-19.71	66.70	15.00
28 PHE CA	-25.93	-19.33	66.91	15.00
28 PHE CB	-26.33	-18.21	65.94	15.00
28 PHE CG	-25.73	-16.88	66.28	15.00
28 PHE CD1	-24.51	-16.49	65.73	15.00
28 PHE CD2	-26.34	-16.04	67.19	15.00
28 PHE CE1	-23.92	-15.30	66.11	15.00
28 PHE CE2	-25.75	-1.4.85	67.57	15.00
28 PHE CZ	-24.54	-14.48	67.02	15.00
28 PHE C	-26.82	-20.55	66.75	15.00
28 PHE O	-27.75	-20.75	67.51	15.00
29 SER N	-26.49	-21.39	65.78	15.00

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29 SER CA	-27.25	-22.61	65.53	15.00
29 SER CB	-26.69	-23.33	64.31	15.00
29 SER OG	-27.48	-24.45	63.98	15.00
29 SER C	-27.16		66.74	15.00
29 SER O	-28.17	-23.93	67.30	15.00
30 SER N	-25.92	-23.82	67.14	15.00
30 SER CA	-25.63	-24.69	68.27	15.00
30 SER CB	-24.13	-24.77	68.46	15.00
30 SER OG	-23.51	-24.96	67.22	15.00
30 SER C	-26.27	-24.16	69.55	15.00
30 SER O	-26.82	-24.92	70.35	15.00
31 VAL N	-26.17	-22.85	69.75	15.00
31 VAL CA	-26.75	-22.24	70.93	15.00
31 VAL CB	-26.25	-20.78	71.09	15.00
31 VAL CG1	-27.32	-19.87	71.63	15.00
31 VAL CG2	-25.06	-20.75	72.01	15.00
31 VAL C	-28.27	-22.35	70.84	15.00
31 VAL O	-28.94	-22.62	71.82	15.00
32 GLY N	-28.80	-22.28	69.63	15.00
32 GLY CA	-30.23	-22.39	69.44	15.00
32 GLY C	-30.80	-23.73	69.86	15.00
32 GLY O	-31.91	-23.80	70.38	15.00
33 ALA N	-30.05	-24.80	69.59	15.00
33 ALA CA	-30.46	-26.15	69.95	15.00
33 ALA CB	-29.65	-27.18	69.17	15.00
33 ALA C	-30.28	-26.35	71.44	15.00
33 ALA O	-31.16	-26.89	72.11	15.00
34 LEU N	-29.14	-25.91	71.97	15.00
34 LEU CA	-2887	-26.01	73.40	15.00
34 LEU CB	-27.54	-25.36	73.74	15.00
34 LEU CG	-26.26	-26.09	73.37	15.00
34 LEU CD1	-25.09	-25.16	73.59	15.00
34 LEU CD2	-26.10	-27.34	74.21	15.00
34 LEU C	-29.98	-25.32	74.18	15.00
34 LEU O	-30.45	-25.85	75.18	15.00
35 GLU N	-30.43	-24.16	73.69	15.00
35 GLU CA	-31.51	-23.39	74.33	15.00
35 GLU CB	-31.65	-22.03	73.66	15.00
35 GLU CG	-30.44	-21.12	73.81	15.00
35 GLU CD	-30.54	~19.82	73.03	15.00
35 GLU OE1	-31.35	-19.71	72.09	15.00
35 GLU OE2	-29.77	-18.91	73.35	15.00
35 GLU C	-32.84	-24.12	74.28	15.00

25 07 11 0				
35 GLU 0	-33.67			15.00
36 GLY N	-33.09		73.21	15.00
36 GLY CA	-34.33		73.09	15.00
36 GLY C	-34.37		74.11	15.00
36 GLY O	-35.37		74.80	15.00
37 GLN N	-33.26		74.25	15.00
37 GLN CA	-33.18		75.20	15.00
37 GLN CB	-31.92		74.98	
37 GLN CG	-31.94		73.69	15.00
37 GLN CD	-33.17		73.57	15.00
37 GLN OE1			74.43	15.00
37 GLN NE2	-33.95	-30.79	72.53	15.00
37 GLN C	-33.24	-28.01	76.63	15.00
37 GLN O	-33.97	-28.56	77.47	15.00
38 LEU N	-32.52	-26.92	76.89	15.00
38 LEU CA	-32.51	-26.31	78.21	15.00
38 LEU CB	-31.79	-24.96	78.17	15.00
38 LEU CG	-31.83	-24.13	79.46	15.00
38 LEU CD1	-31.00	-24.83	80.53	15.00
38 LEU CD2	-31.30	-22.73	79.20	15.00
38 LEU C	-33.94	-26.11	78.69	15.00
38 LEU O	-34.30	-26.51	79.79	15.00
39 LYS N	-34.77	-25.51	77.85	15.00
39 LYS CA	-36.16	-25.26	78.20	15.00
39 LYS CB	-36.85	-24.43	77.11	15.00
39 LYS CG	-38.06	-23.64	77.59	15.00
39 LYS CD	-39.32	-24.48	77.65	15.00
39 LYS CE	-40.53	-23.66	78.07	15.00
39 LYS NZ	-40.50	-23.27	79.50	15.00
39 LYS C	-36.89		78.36	15.00
39 LYS O	-37.71	-26.75	79.25	15.00
40 LYS N	-36.57	-27.57	77.53	15.00
40 LYS CA	-37.25	-28.86	77.63	15.00
40 LYS CB	-36.85		76.48	15.00
40 LYS CG	-37.77	-31.00	76.39	15.00
40 LYS CD	-37.48	-31.90	75.22	
40 LYS CE	-38.53	-32.99		15.00
40 LYS NZ	-38.45	-33.78	75.17	15.00
40 LYS C		-33.78	73.92	15.00
40 LYS O		-30.11	78.95	15.00
41 LYS N			79.56	15.00
41 LYS CA		-29.45	79.42	15.00
41 LYS CB		-30.09	80.66	15.00
225 CB	~33.8/	-30.48	80.61	15.00

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41 LYS CG	-33.50		79.43	15.00
41 LYS CD	-34.48		79.32	
41 LYS CE	-34.33	_		
41 LYS NZ	-35.59			
41 LYS C	-35.67		81.94	
41 LYS O	-36.36		82.81	15.00
42 THR N	~35.20		82.07	
42 THR CA	-35.45		83.29	15.00
42 THR CB	-34.31		83.57	15.00
42 THR OG1	-34.40		82.67	15.00
42 THR CG2	-32.98	-27.04	83.36	15.00
42 THR C	-36.77		83.35	15.00
42 THR O	-37.19		84.43	15.00
43 GLY N	-37.42		82.21	15.00
43 GLY CA	-38.67		82.16	15.00
43 GLY C	-38.49	-24.09	82.05	15.00
43 GLY O	-39.45	-23.33	81.92	15.00
44 LYS N	-37.25	-23.62	82.11	15.00
44 LYS CA	-36.99		82.03	15.00
44 LYS CB	-36.47	-21.70	83.37	15.00
44 LYS CG	-37.07	-22.42	84.57	15.00
44 LYS CD	-36.69	-21.78	85.90	15.00
44 LYS CE	-35.51	-22.46	86.59	15.00
44 LYS NZ	-35.95	-23.54	87.52	15.00
44 LYS C	-35.98	-21.94	80.92	15.00
44 LYS 0	-34.92	-22.58	80.87	15.00
45 LEU N	-36.32	-21.02	80.04	15.00
45 LEU CA 45 LEU CB	-35.47	-20.64	78.90	15.00
	-36.33	-20.13	77.75	15.00
45 LEU CG 45 LEU CD1	-35.67	-19.77	76.42	15.00
45 LEU CD1 45 LEU CD2	-35.27	-21.03	75.66	15.00
45 LEU CD2	-36.67	-18.97	75.62	15.00
45 LEU O	-34.47	-19.58	79.29	15.00
	-34.68	-18.81	80.22	15.00
	-33.38	-19.50	78.54	15.00
	-32.33	-18.54	78.81	15.00
46 LEU CB 46 LEU CG	-31.43	-19.09	79.91	15.00
	-30.45	-18.17	80.63	15.00
	-31.17	-16.89	81.03	15.00
	-29.91	-18.88	81.84	15.00
46 LEU C	-31.55	-18 33	77.51	15.00
16 LEU O	-31.62	-19.14	76.60	15.00
17 ASN N	-30.84	-17.22	77.39	15.00

47 ASN CA	-30.04	-16.97	76.20	15.00
47 ASN CB	-30.04		75.85	15.00
47 ASN CG	-31.37		75.33	15.00
47 ASN OD1	-32.19	-14.49	76.08	15.00
47 ASN ND2	-31.59	-15.20	74.05	15.00
47 ASN C	-28.62	-17.42	76.46	15.00
47 ASN O	-27.96		77.37	15.00
48 LEU N	-28.15	-18.40	75.69	15.00
48 LEU CA	-26.80	-18.91	75.86	15.00
48 LEU CB	-26.74	-20.39	75.53	15.00
48 LEU CG	-27.64	-21.29	76.40	15.00
48 LEU CD1	-27.37		76.07	15.00
48 LEU CD2	-27.39		77.86	15.00
48 LEU C	-25.79	-18.10	75.06	15.00
48 LEU O	-26.16	-17.33	74.17	15.00
49 SER N	-24.51	-18.27	75.36	15.00
49 SER CA	-23.45	-17.50	74.71	15.00
49 SER CB	-22.34	-17.20	75.73	15.00
49 SER OG	-21.21	-16.57	75.14	15.00
49 SER C	-22.81	-18.05	73.44	15.00
49 SER 0	-22.00	-18.98	73.49	15.00
50 PRO N	-23.17	-17.49	72.28	15.00
50 PRO CD	-24.25	-16.53	71.99	15.00
50 PRO CA	-22.56	-17.98	71.04	15.00
50 PRO CB	-23.40	-17.29	69.95	15.00
50 PRO CG	-23.89	-16.05	70.62	15.00
50 PRO C	-21.10	-17.55	71.00	15.00
50 PRO O	-20.25	-18.21	70.41	15.00
51 GLN N	-20.79	-16.45	71.69	15.00
51 GLN CA	-19.43	-15.93	71.75	15.00
51 GLN CB	-19.40	-14.60	72.52	15.00
51 GLN CG	-18.07		72.44	15.00
51 GLN CD	-17.86	-13.13	71.13	15.00
51 GLN OE1	-18.70	-12.34	70.70	15.00
51 GLN NE2	-16.72		70.49	15.00
51 GLN C	-18.52		72.44	15.00
51 GLN O	-17.43		71.95	15.00
52 ASN N	-18.97	-17.50	73.56	15.00
52 ASN CA		-18.49	74.31	15.00
52 ASN CB	-19.13			15.00
52 ASN CG		-20.00		15.00
52 ASN OD1				15.00
52 ASN ND2	-17.40	-20.73	75.93	15.00

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52 ASN C	-17.77		73.34	15.0
52 ASN 0	-16.66		73.43	
53 LEU N	-18.65		72.39	15.0
53 LEU CA	-18.37		71.39	
53 LEU CB	-19.67		70.73	15.00
53 LEU CG	-20.66	-22.13	71.63	15.00
53 LEU CD1	-21.88	-22.54	70.82	
53 LEU CD2	-20.00	-23.35	72.23	-
53 LEU C	-17.40		70.35	15.00
53 LEU O	-16.45		69.98	15.00
54 VAL N	-17.59		69.91	15.00
54 VAL CA	-16.73		68.89	15.00
54 VAL CB	-17.24		68.51	15.00
54 VAL CG1	-16.31		67.49	15.00
54 VAL CG2	-18.63	-17.19	67.92	15.00
54 VAL C	-15.27		69.36	15.00
54 VAL O	-14.36		68.64	15.00
55 ASP N	-15.06		70.58	15.00
55 ASP CA	-13.73		71.15	15.00
55 ASP CB	-13.78		72.29	15.00
55 ASP CG	-14.32		71.87	15.00
55 ASP OD1	-14.20	-15.10	70.68	15.00
55 ASP OD2	-14.86	-14.76	72.75	15.00
55 ASP C	-13.05	-19.09	71.70	15.00
55 ASP 0	-11.81	-19.13	71.78	15.00
56 CYS N	-13.81	-20.12	72.07	15.00
56 CYS CA	-13.22	-21.32	72.67	15.00
56 CYS C	-13.29	-22.63	71.91	15.00
56 CYS O	-12.56	-23.59	72.23	15.00
56 CYS CB	-13.83	-21.53	74.05	15.00
56 CYS SG	-14.01	-20.02	75.04	15.00
57 VAL N	-14.19	-22.77	70.96	15.00
57 VAL CA	-14.26	-24.02	70.22	15.00
57 VAL CB	-15.60	-24.16	69.48	15.00
57 VAL CG1	-15.62	-25.44	68.67	15.00
57 VAL CG2	-16.73	-24.15	70.48	15.00
57 VAL C	-13.09	-24.06	69.25	
57 VAL O	-13.23	-23.80	68.06	15.00
58 SER N	-11.91	-24.40	69.77	15.00
58 SER CA		-24.46	69.00	15.00
58 SER CB		-24.83	69.93	15.00
58 SER OG		-2 <b>4</b> .83		15.00
58 SER C		-25.38	70.98 67.79	15.00
		43.30	01.19	15.00

58 SER 0	-9.70	-25.38	67.02	15.00
59 GLU N	-11.70	-26.18	67.64	15.00
59 GLU CA	-11.82	-27.10	66.50	15.00
59 GLU CB	-12.70	-28.31	66.86	15.00
59 GLU CG	-12.16	-29.19	67.99	15.00
59 GLU CD	-12.31	-28.56	69.36	15.00
59 GLU OE1	-13.46	-28.38	69.83	15.00
59 GLU OE2	-11.28	-28.23	69.97	15.00
59 GLU C	-12.44	-26.34	65.33	15.00
59 GLU O	-12.44	-26.81	64.19	15.00
60 ASN N	-13.03	-25.19	65.61	15.00
60 ASN CA	-13.64	-24.37	64.58	15.00
60 ASN CB	-15.08	-23.98	64.97	15.00
60 ASN CG	-16.03	-25.16	64.95	15.00
60 ASN OD1	-17.08	-25.12	65.58	15.00
60 ASN ND2	-15.68	-26.20	64.22	15.00
60 ASN C	-12.79	-23.12	64.31	15.00
60 ASN O	-11.88	-22.80	65.06	15.00
61 ASP N	-13.13	-22.40	63.25	15.00
61 ASP CA	-12.38	-21.21	62.87	15.00
61 ASP CB	-12.30	-21.14	61.34	15.00
61 ASP CG	-10.95	-20.64	60.84	15.00
61 ASP OD1	-10.04	-20.40	61.65	15.00
61 ASP OD2	-10.80	-20.50	59.60	15.00
61 ASP C	-12.97	-19.91	63.44	15.00
61 ASP O	-12.70	-18.83	62.93	15.00
62 GLY N	-13.78	-19.99	64.49	15.00
62 GLY CA	-14.37	-18.79	65.05	15.00
62 GLY C	-15.25	-18.07	64.05	15.00
62 GLY 0	-16.26	-18.59	63.58	15.00
63 CYS N	-14.89	-16.85	63.70	15.00
63 CYS CA	-15.67	-16.11	62.72	15.00
63 CYS C	-15.48	-16.70	61.33	15.00
63 CYS O	-16.15		60.39	15.00
63 CYS CB	-15.30		62.69	15.00
63 CYS SG		-13.63	64.12	15.00
64 GLY N	-14.54	-17.62	61.19	15.00
64 GLY CA	-14.29	-18.25	59.90	15.00
64 GLY C		-19.41	59.67	15.00
64 GLY O		-19.97	58.57	15.00
65 GLY N		-19.80	60.70	15.00
65 GLY CA		-20.91	60.56	15.00
65 GLY C	-16.55	-22.16	61.33	15.00

65 GLY O	-15.46	-22.30	61.90	15.00
66 GLY N	-17.47	-23.10	61.38	15.00
66 GLY CA	-17.25	-24.34	62.07	15.00
66 GLY C	-18.51	-25.15	61.93	15.00
66 GLY O	-19.46	-24.72	61.29	15.00
67 TYR N	-18.50	-26.35	62.49	15.00
67 TYR CA	-19.66	-27.22	62.44	15.00
67 TYR CB	-19.25	-28.66	62.13	15.00
67 TYR CG	-18.41	-28.83	60.89	15.00
67 TYR CD	L -19.00	-29.00	59.64	15.00
67 TYR CE	L -18.22	-29.18	58.51	15.00
67 TYR CD2	-17.03	-28.83	60.98	15.00
67 TYR CE2	-16.25	-29.01	59.86	15.00
67 TYR CZ	-16.84	-29.18	58.63	15.00
67 TYR OH	-16.03	-29.34	57.52	15.00
67 TYR C	-20.27	-27.16	63.83	15.00
67 TYR O	-19.59	-26.85	64.80	15.00
68 MET N	-21.55	-27.48	63.93	15.00
68 MET CA	-22.24	-27.45	65.20	15.00
68 MET CB	-23.75	-27.51	65.00	15.00
68 MET CG	-24.34	-26.31	64.22	15.00
68 MET SD	-23.95	-26.25	62.46	15.00
68 MET CE	-25.40	-26.96	61.77	15.00
68 MET C	-21.76	-28.58	66.11	15.00
68 MET O	-21.49	-28.37	67.29	15.00
69 THR N	-21.57	-29.78	65.56	15.00
69 THR CA	-21.13	-30.94	66.33	15.00
69 THR CB	-20.92	-32.18	65.44	15.00
69 THR OG1	-20.10	-31.83	64.31	15.00
69 THR CG2	-22.26	-32.69	64.93	15.00
69 THR C	-19.85	-30.65	67.12	15.00
69 THR O	-19.69	-31.11	68.26	15.00
70 ASN N	-18.95	-29.84	66.55	15.00
70 ASN CA	-17.71	-29.46	67.22	15.00
70 ASN CB	-16.73		66.24	15.00
70 ASN CG	-15.97	-29.81	65.39	15.00
70 ASN OD1	-15.41	-29.45	64.37	15.00
70 ASN ND2	-15.92	-31.06	65.83	15.00
70 ASN C.	-17.96	-28.52	68.39	15.00
70 ASN O	-17.14	-28.42	69.30	15.00
71 ALA N		-27.78	68.35	15.00
71 ALA CA	-19.42	-26.86	69.44	15.00
71 ALA CB	-20.32	-25.76	68.92	15.00

71 ALA C	-20.12	-27.66	70.53	15 00
71 ALA 0	-19.94			
72 PHE N	-20.96		70.13	15.00
72 PHE CA	-21.67		71.08	
72 PHE CB	-22.56		70.34	15.00
72 PHE CG	-23.74		69.63	15.00
72 PHE CD1	-24.33		70.11	
72 PHE CD2	-24.27		68.50	
72 PHE CE1	-25.45		69.48	
72 PHE CE2	-25.39		67.87	15.00
72 PHE CZ	-25.98		68.35	15.00
72 PHE C	-20.60		71.89	15.00
72 PHE 0	-20.60		73.12	15.00
73 GLN N	-19.64		71.21	15.00
73 GLN CA	-18.58	-31.56	71.87	15.00
73 GLN CB	-17.64	-32.18	70.82	15.00
73 GLN CG	-16.55		71.36	15.00
73 GLN CD	-17.07		71.69	15.00
73 GLN 0E1	-17.16		72.87	15.00
73 GLN NE2	-17.37	-35.32	70.65	15.00
73 GLN C	-17.81	-30.69	72.87	15.00
73 GLN O	-17.46		73.96	15.00
74 TYR N	-17.55		72.53	15.00
74 TYR CA	-16.82	-28.56	73.44	15.00
74 TYR CB	-16.43	-27.25	72.75	15.00
74 TYR CG	-16.11	-26.12	73.70	15.00
74 TYR CD1	-14.83	-25.93	74.20	15.00
74 TYR CE1	-14.54	-24.93	75.11	15.00
74 TYR CD2	-17.11	-25.25	74.14	15.00
74 TYR CE2	-16.84	-24.23	75.06	15.00
74 TYR CZ	-15.55	-24.08	75.53	15.00
74 TYR OH	-15.28		76.45	15.00
74 TYR C	-17.62		74.70	15.00
74 TYR O	-17.05		75.76	15.00
75 VAL N	-18.94		74.59	15.00
75 VAL CA		-27.89	75. <b>7</b> 5	15.00
75 VAL CB		-27.54	75.33	15.00
75 VAL CG1	-22.04	-27.14	76.54	15.00
75 VAL CG2	-21.20	-26.39	74.33	15.00
75 VAL C		-29.11	76.68	15.00
75 VAL 0		-28.99	77.90	15.00
76 GLN N	-19.50		76.09	15.00
76 GLN CA	-19.43	-31.54	76.84	15.00

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76 GLN CB	-19.51	-32.72	75.88	15.00
76 GLN CG	-19.44	-34.10		15.00
76 GLN CD	-19.41	-35.21	75.50	15.00
76 GLN OE1	-20.45	-35.67	75.05	15.00
76 GLN NE2	-18.23	-35.64	75.12	15.00
76 GLN C	-18.15	-31.64	77.66	15.00
76 GLN 0	-18.20	-31.71	78.89	15.00
77 LYS N	-17.00	-31.64	77.00	15.00
77 LYS CA	-15.72	-31.75	77.70	15.00
77 LYS CB	-14.60	-32.13	76.73	15.00
77 LYS CG	-14.72	-31.53	75.36	15.00
77 LYS CD	-13.52	-31.92	74.50	15.00
77 LYS CE	-13.77	-31.61	73.03	15.00
77 LYS NZ	-14.25	-30.21	72.83	15.00
77 LYS C	-15.29	-30.55	78.56	15.00
77 LYS O	-14.36	-30.63	79.36	15.00
78 ASN N	-15.94	-29.41	78.38	15.00
78 ASN CA	-15.62	-28.22	79.15	15.00
78 ASN CB	-15.75	-26.98	78.27	15.00
78 ASN CG	-15.54	-25.69	79.03	15.00
78 ASN OD1	-14.43	-25.40	79.46	15.00
78 ASN ND2	-16.60	-24.91	79.16	15.00
78 ASN C	-16.60	-28.14	80.32	15.00
78 ASN O	-16.47	-27.32	81.22	15.00
79 ARG N	-17.60	-29.01	80.27	15.00
79 ARG CA	-18.63	-29.08	81.29	15.00
79 ARG CB	-18.06	-29.64	82.60	15.00
79 ARG CG	-17.45	-31.04	82.42	15.00
79 ARG CD	-17.13	-31.71	83.75	15.00
79 ARG NE	-18.34	-32.20	84.40	15.00
79 ARG CZ	-18.91	-33.37	84.15	15.00
79 ARG NH1	-18.36	-34.20	83.26	15.00
79 ARG NH2		-33.70	84.75	15.00
79 ARG C	-19.36	-27.77	81.50	15.00
79 ARG O		-27.33	82.64	15.00
80 GLY N	-19.75	-27.14	80.39	15.00
80 GLY CA	-20.48	-25.89	80.49	15.00
80 GLY C	-20.34	-24.84	79.40	15.00
80 GLY O	-19.34	-24.78	78.67	15.00
81 ILE N	-21.39	-24.02	79.30	15.00
81 ILE CA	-21.46	-22.91	78.36	15.00
81 ILE CB	-22.22	-23.28	77.05	15.00
81 ILE CG2	-23.65	-23.63	77.33	15.00
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81 ILE CG1	-22 16	-22.09	25.00	
81 ILE CD1	-22.80			
81 ILE C	-22.15		_	15.00
81 ILE 0	-23.20	_	79.07	15.00
82 ASP N	-21.51		79.71	15.00
82 ASP CA	-22.01		79.03	
82 ASP CB	-20.99			
82 ASP CG	-19.78		79.46	
82 ASP OD1	-18.71		80.35	
82 ASP OD2	-19.91		79.95	
82 ASP C	-23.36		81.47	15.00
82 ASP O		-10.86	79.13	15.00
83 SER N	-23.91		78.09	15.00
83 SER CA	-25.16		79.86	15.00
83 SER CB	-25.99		79.50	15.00
83 SER OG	-25.21		80.76	
83 SER C	-24.75		81.78	15.00
83 SER O	-23.63		78.85	15.00
84 GLU N	-25.64	-15.45	79.05	15.00
84 GLU CA	-25.32	-14.09	78.07	15.00
84 GLU CB	-26.48		77.39	15.00
84 GLU CG	-26.16	-13.65	76.52	15.00
84 GLU CD	-25.34	-12.95	75.57	15.00
84 GLU OE1	-24.85	-14.10	74.38	15.00
84 GLU OE2	-25.19	-12.13	74.35	15.00
84 GLU C	-24.88		73.45	15.00
84 GLU O	-24.03	-12.14	78.31 77.94	15.00
85 ASP N	-25.44	-12.87	79.52	15.00
85 ASP CA	-25.03	-11.81	80.45	15.00
85 ASP CB	-25.92		81.71	15.00
85 ASP CG	-25.48		82.73	15.00 15.00
85 ASP OD1	-24.60	-9.85	82.41	15.00
85 ASP OD2	-26.00	-10.66	83.87	15.00
85 ASP C		-12.02	80.86	15.00
85 ASP O	-22.83		81.00	15.00
86 ALA N	-23.18		81.05	15.00
86 ALA CA	-21.82	-13.56	81.47	15.00
86 ALA CB	-21.76	-14.91	82.14	15.00
86 ALA C	-20.78	-13.48	80.36	15.00
86 ALA O	-19.62	-13.15	80.59	15.00
87 TYR N		-13.74	79.12	15.00
87 TYR CA		-13.71	78.02	15.00
87 TYR CB	-19.79	-15.15	78.02	15.00
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87		-18.44	-15.33	77.08	15.00
87		-17.91	-14.37	76.21	15.00
87		-16.69	-14.59	75.57	15.00
87	TYR CD2	-17.72	-16.51	77.28	15.00
87	TYR CE2	-16.50	-16.73	76.64	15.00
87		-16.00	-15.77	75.79	15.00
87	TYR OH	-14.79	-16.02	75.16	15.00
87	TYR C	-20.92	-13.13	76.79	15.00
87	TYR O	-21.26	-13.88	75.86	15.00
88	PRO N	-21.19	-11.80	76.79	15.00
88	PRO CD	-21.02	-10.93	77.96	15.00
88	PRO CA	-21.85	-11.07	75.70	15.00
88	PRO CB	-21.91	-9.64	76.23	15.00
88	PRO CG	-22.06	-9.86	77.72	15.00
88	PRO C	-21.13	-11.13	74.36	15.00
88	PRO O	-19.90	-11.11	74.31	15.00
89	TYR N	-21.93	-11.18	73.31	15.00
89	TYR CA	-21.47	-11.28	71.92	15.00
89	TYR CB	-22.64	-11.80	71.09	15.00
89	TYR CG	-22.28	-12.14	69.68	15.00
89	TYR CD1	-21.43	-13.20	69.39	15.00
89	TYR CE1	-21.07	-13.48	68.09	15.00
89	TYR CD2	-22.75	-11.36	68.63	15.00
89	TYR CE2	-22.39	-11.63	67.32	15.00
89	TYR CZ	-21.55	-12.69	67.06	15.00
89	TYR OH	-21.15	-12.93	<b>65.7</b> 7	15.00
89	TYR C	-20.94	-9.96	71.34	15.00
89	TYR O	-21.71	-9.04	71.08	15.00
90	VAL N	-19.64	-9.86	71.12	15.00
90	VAL CA	-19.06	-8.63	70.58	15.00
90	VAL CB	-17.75	-8.27	71.27	15.00
90	VAL CG1	-17.98	-8.12	72.76	15.00
90	VAL CG2	-16.69	-9.32	70.99	15.00
	VAL C	-18.87	-8.65	69.07	15.00
	VAL O	-18.28	-7.74	68.50	15.00
91		-19.30	-9.73	68.44	15.00
91		-19.20	-9.84	67.00	15.00
91	GLY C	-17.86	-10.11	66.36	15.00
91	GLY O	-17.67	-9.84	65.18	15.00
92	GLN N	-16.91	-10.66	67.11	15.00
92	GLN CA	-15.59	-10.97	66.57	15.00
92		-14.80	-9.69	66.32	15.00
92	GLN CG	-14.73	-8.76	67.51	15.00

92 GLN CD	-13.95	-7.50	67.19	15.00
92 GLN OE1	-12.73	-7.54	67.06	15.00
92 GLN NE2	-14.65	-6.38	67.02	15.00
92 GLN C	-14.83	-11.87	67.53	15.00
92 GLN O	-15.17	-11.94	68.71	15.00
93 GLU N	-13.80	-12.53	67.04	15.00
93 GLU CA	-13.02	-13.45	67.85	15.00
93 GLU CB	-12.36	-14.52	66.98	15.00
93 GLU CG	-11.54	-13.97	65.80	15.00
93 GLU CD	-11.57	-14.90	64.59	15.00
93 GLU OE1	-10.51	-15.16	63.98	15.00
93 GLU OE2	-12.67	-15.36	64.22	15.00
93 GLU C	-11.99	-12.78	68.74	15.00
93 GLU 0	-11.29	-11.86	68.33	15.00
94 GLU N	-11.89	-13.29	69.97	15.00
94 GLU CA	-10.96	-12.78	70.96	15.00
94 GLU CB	-11.65	-11.74	71.84	15.00
94 GLU CG	-12.68	-12.34	72.79	15.00
94 GLU CD	-13.96	-11.54	72.88	15.00
94 GLU OE1	-13.96	-10.36	72.44	15.00
94 GLU OE2	-14.97	-12.08	73.38	15.00
94 GLU C	-10.54	-13.98	71.82	15.00
94 GLU 0	-10.95	-15.10	71.55	15.00
95 SER N	-9.72	-13.74	72.84	15.00
95 SER CA	-9.26	-14.80	73.73	15.00
95 SER CB	-8.26	-14.23	74.75	15.00
95 SER OG	-7.14	-13.65	74.09	15.00
95 SER C	-10.41	-15.51	74.46	15.00
95 SER 0	-11.34	-14.87	74.95	15.00
96 CYS N	-10.37	-16.84	74.53	15.00
96 CYS CA	-11.44	-17.57	75.21	15.00
96 CYS C	-11.55	-17.03	76.62	15.00
96 CYS O	-10.58	-17.04	77.37	15.00
96 CYS CB		-19.06	75.26	15.00
96 CYS SG 97 MET N	-12.35	-20.02	76.20	15.00
	-12.72	-16.51	76.97	15.00
	-12.95	-15.95	78.31	15.00
7 MET CB	-13.50	-14.53	78.21	15.00
7 MET CG 7 MET SD	-12.65	-13.51	77.52	15.00
	-13.49	-11.94	77.83	15.00
7 MET CE	-14.95	-12.05	76.75	15.00
7 MET C	-13.95	-16.74	79.15	15.00
7 MET O	-14.60	-16.15	80.02	15.00

98 TYR N	-14.08	-18.05	78.94	15.00
98 TYR CA	-15.05		79.70	15.00
98 TYR CB	-14.89		79.47	
98 TYR CG	-15.95	-21.14	80.19	15.00
98 TYR CD1	-17.30	-21.01	79.88	15.00
98 TYR CE1	-18.28	-21.71	80.58	15.00
98 TYR CD2	-15.61	-22.00	81.23	15.00
98 TYR CE2	-16.58		81.94	15.00
98 TYR CZ	-17.91	_	81.62	15.00
98 TYR OH	-18.86		82.33	15.00
98 TYR C	-14.99		81.20	15.00 15.00
98 TYR O	-13.92		81.77	15.00
99 ASN N	-16.16		81.82	15.00
99 ASN CA	-16.30		83.23	15.00
99 ASN CB	-16.73	-16.72	83.37	15.00
99 ASN CG	-17.06	-16.32	84.81	15.00
99 ASN OD1	-16.96		85.74	15.00
99 ASN ND2	-17.47		84.98	15.00
99 ASN C	-17.34	-19.10	83.86	15.00
99 ASN 0	-18.55	-18.87	83.72	15.00
100 PRO N	-16.89	-20.13	84.60	15.00
100 PRO CD	-15.46	-20.39	84.86	15.00
100 PRO CA	-17.73	-21.12	85.29	15.00
100 PRO CB	-16.74	-21.80	86.22	15.00
100 PRO CG	-15.49	-21.81	85.41	15.00
100 PRO C	-18.84	-20.46	86.07	15.00
100 PRO O	-19.93	-21.02	86.20	15.00
101 THR N	-18.58	-19.26	86.58	15.00
101 THR CA	-19.58	-18.53	87.34	15.00
101 THR CB	-18.99	-17.20	87.84	15.00
101 THR OG1	-17.76	-17.48	88.53	15.00
101 THR CG2	-19.95	-16.50	88.80	15.00
101 THR C	-20.78	-18.25	86.46	15.00
101 THR 0	-21.93	-18.29	86.93	15.00
102 GLY N	-20.52	-17.99	85.18	15.00
102 GLY CA	-21.57	-17.69	84.23	15.00
102 GLY C	-22.29	-18.89	83.62	15.00
102 GLY 0	-23.36	-18.73	83.04	15.00
103 LYS N	-21.70	-20.07	83.72	15.00
103 LYS CA	-22.30	-21.29	83.17	15.00
103 LYS CB		-22.52	83.76	15.00
103 LYS CG	-22.34	-23.84	83.55	15.00
103 LYS CD	-21.59	-34.99	84.23	15.00

103	LYS	CE	-22.56	-26.02	84.79	15.00
103	LYS	NZ	-23.48	-26.58	83.76	15.00
103	LYS	С	-23.81	-21.37	83.41	15.00
103	LYS	0	-24.27	-21.24	84.54	15.00
104	ALA	N	-24.58	-21.60	82.35	15.00
104	ALA	CA	-26.04	-21.70	82.45	15.00
104	ALA	CB	-26.69	-20.52	81.78	15.00
104	ALA	С	-26.60	-22.99	81.88	15.00
104	ALA	0	-27.76	-23.32	82.12	15.00
105	ALA	N	-25.81	-23.71	81.09	15.00
105	ALA	CA	-26.26	-24.97	80.50	15.00
105	ALA	CB	-26.99	-24.72	79.19	15.00
105	ALA	C	-25.10	-25.92	80.28	15.00
105	ALA	0	-23.93	-25.52	80.38	15.00
106	LYS	N	-25.41	-27.18	79.98	15.00
106	LYS	CA	-24.40	-28.19	79.72	15.00
106	LYS	CB	-23.77	-28.66	81.03	15.00
	LYS		-24.73	-29.25	82.04	15.00
	LYS		-24.01	-29.56	83.33	15.00
	LYS	CE	-22.75	-30.37	83.07	15.00
	LYS		-23.05	-31.63	82.31	15.00
	LYS	С	-25.06	-29.34	78.99	15.00
	LYS	0	-26.28	-29.39	78.94	15.00
107	CYS		-24.27	-30.21	78.38	15.00
	CYS		-24.83	-31.36	77.65	15.00
	CYS		-25.14	-30.97	76.20	15.00
	CYS		-23.71	-31.01	75.11	15.00
		С	-23.95	-32.61	77.67	15.00
			-22.73	-32.53	77.81	15.00
	ARG		-24.59	-33.76	77.53	15.00
	ARG		-23.92	-35.05	77.54	15.00
	ARG		-24.66	-36.03	78.46	15.00
	ARG		-26.18	-35.81	78.55	15.00
	ARG			-37.01	79.17	15.00
	ARG		-27.06		78.25	15.00
	ARG			-38.49	77.62	15.00
	ARG			-37.79	77.80	15.00
	ARG		-28.22	-39.56	76.84	15.00
	ARG		-23.70	-35.67	76.15	15.00
	ARG			-36.88	75.98	15.00
	GLY :			-34.83	75.16	15.00
	GLY			-35.32	73.82	
109	GLY ·	C	-24.08	-34.73	72.75	15.00

109	GLY	0	-24.72	-33.69	72.94	15.00
110	TYR	N	-24.15	-35.41	71.61	15.00
110	TYR	CA	-24.97	-34.95	70.50	15.00
110	TYR	CB	-24.19	-33.91	69.71	15.00
110	TYR	CG	-22.97	-34.46	68.99	15.00
110	TYR	CD1	-21.71	-34.43	69.59	15.00
110	TYR	CE1	-20.59	-34.90	68.91	15.00
110	TYR	CD2	-23.07	-34.99	67.71	15.00
110	TYR	CE2	-21.97	-35.45	67.03	15.00
110	TYR	CZ	-20.73	-35.40	67.63	15.00
110	TYR	OH	-19.63	-35.82	66.93	15.00
110	TYR	С	-25.31	-36.11	69.57	15.00
110	TYR	0	-24.61	-37.12	69.54	15.00
111	ARG	N	-26.35	-35.94	68.76	15.00
111	ARG	CA	-26.74	-36.97	67.82	15.00
111	ARG	CB	-28.02	-37.68	68.29	15.00
111	ARG	CG	-27.87	-38.44	69.61	15.00
111	ARG	CD	-29.17	-39.16	70.00	15.00
111	ARG	NE	-29.49	-40.25	69.07	15.00
111	ARG	CZ	-28.84	-41.41	69.03	15.00
111	ARG	NH1	-29.20	-42.34	68.14	15.00
111	ARG	NH2	-27.85	-41.65	69.88	15.00
111	ARG	C	-27.00	-36.34	66.45	15.00
111	ARG	0	-27.71	-35.34	66.34	15.00
112	GLU	N	-26.39	-36.90	65.42	15.00
112	GLU	CA	-26.57	-36.42	64.06	15.00
112	GLU	CB	-25.27	-36.56	63.27	15.00
112	GLU	CG	-24.17	-35.62	63.74	15.00
112	GLU	CD	-22.80	-35.91	63.12	15.00
112	GLU	OE1	-22.22	-34.99	62.47	15.00
112	GLU	OE2	-22.29	-37.05	63.30	15.00
112	GLU	С	-27.68	-37.26	63.45	15.00
112	GLU	0	-27.84	-38.43	63.80	15.00
113	ILE	N	-28.50	-36.63	62.61	15.00
113	ILE	CA	-29.60	-37.28	61.90	15.00
113	ILE	CB	-30.69	-36.23	61.48	15.00
113	ILE	CG2	-31.45	-36.68	60.25	15.00
113	ILE	CG1	-31.68	-35.99	62.62	15.00
113	ILE	CD1	-31.14	-35.16	63.73	15.00
113	ILE	C	-29.00	-37.94	60.66	15.00
113	ILE	0	-28.03	-37.44	60.10	15.00
114	PRO	N	-29.54	-39.09	60.23	15.00
114	PRO	CD	-30.62	-39.89	60.83	15.00

114	PRO	CA	-29.00	-39.75	59.04	15.00
114	PRO	CB	-30.00	-40.89	58.81	15.00
114	PRO	CG	-30.41	-41.24	60.19	15.00
114	PRO	C	-28.97	-38.80	57.86	15.00
114	PRO	0	-29.98	-38.20	57.52	15.00
115	GLU	N	-27.80	-38.73	57.22	15.00
115	GLU	CA	-27.56	-37.86	56.07	15.00
115	GLU	CB	-26.14	-38.07	55.52	15.00
115	GLU	CG	-25.92	-37.52	54.10	15.00
115	GLU	CD	-24.48	-37.06	53.82	15.00
115	GLU	OE1	-23.52	-37.73	54.28	15.00
115	GLU	OE2	-24.30	-36.02	53.15	15.00
115	GLU	С	-28.55	-38.00	54.93	15.00
115	GLU	0	-28.57	-39.02	54.26	15.00
116	GLY	N	-29.31	-36.94	54.69	15.00
116	GLY	CA	-30.27	-36.94	53.60	15.00
116	GLY	С	-31.66	-37.41	53.97	15.00
116	GLY	0	-32.57	-37.40	53.14	15.00
117	asn	N	-31.86	-37.78	55.24	15.00
117	ASN	CA	-33.15	-38.29	55.69	15.00
117	ASN	CB	-32.94	-39.38	56.73	15.00
117	asn	CG	-34.19	-40.18	56.99	15.00
117	ASN	OD1	-35.32	-39.68	56.89	15.00
	ASN		-34.00	-41.46	57.33	15.00
	ASN		-34.07	-37.22	56.27	15.00
	asn		-33.93	-36.82	57.43	15.00
118	GLU	N	-35.06	-36.79	55.50	15.00
118	GLU	CA	-35.97	-35.78	55.99	15.00
118	GLU	CB	-36.70	-35.07	54.85	15.00
	GLU		-36.04	-33.77	54.42	15.00
	GLU	CD	-36.91	-32.99	53.45	15.00
			-36.81	-33.22	52.23	15.00
	GLU		-37.72	-32.15	53.91	15.00
	GLU		-36.97		56.98	15.00
	GLU		-37.46	-35.59	57.83	15.00
	LYS				56.90	15.00
	LYS		-38.27	-38.14	57.87	15.00
	LYS			-39.49	57.42	15.00
		CG		-39.34	56.68	15.00
119				-38.40	55.47	15.00
	LYS			-38.06		15.00
	LYS			-37.19		15.00
119	LYS	С	-37.62	-38.24	59.24	15.00

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119 LYS O	-38.23	-37.90	60.26	15.00
120 ALA N	-36.35		59.28	15.00
120 ALA CA	-35.63	· · · · · ·	60.54	15.00
120 ALA CB	-34.25	=	60.33	15.00
120 ALA C	-35.55	-37.30	61.16	15.00
120 ALA O	-35.58	-37.15	62.38	15.00
121 LEU N	-35.45	-36.27	60.32	15.00
121 LEU CA	-35.38	-34.89	60.82	15.00
121 LEU CB	-34.93	-33.91	59.73	15.00
121 LEU CG	-34.80	-32.45	60.19	15.00
121 LEU CD1	-33.77	-32.34	61.30	15.00
121 LEU CD2	-34.42	-31.56	59.03	15.00
121 LEU C	-36.74	-34.48	61.36	15.00
121 LEU 0	-36.83	-33.80	62.38	15.00
122 LYS N	-37.79	-34.89	60.66	15.00
122 LYS CA	-39.16	-34.58	61.05	15.00
122 LYS CB	-40.14	-35.19	60.07	15.00
122 LYS CG	-41.57	-34.80	60.31	15.00
122 LYS CD	-42.49	-35.45	59.32	15.00
122 LYS CE	-43.95	-35.16	59.67	15.00
122 LYS NZ	-44.88	-35.66	58.60	15.00
122 LYS C	-39.41	-35.15	62.44	15.00
122 LYS 0	-39.87	-34.44	63.33	15.00
123 ARG N	-39.12	-36.44	62.62	15.00
123 ARG CA	-39.29	-37.10	63.90	15.00
123 ARG CB	-38.84	-38.58	63.84	15.00
123 ARG CG	-39.74	-39.50	62.99	15.00
123 ARG CD	-39.33	-40.97	63.13	15.00
123 ARG NE	-37.95	-41.22	62.73	15.00
123 ARG CZ	-37.42	-42.43	62.51	15.00
123 ARG NH1	-38.15	-43.53	62.66	15.00
123 ARG NH2	-36.17	-42.53	62.08	15.00
123 ARG C	-38.46	-36.37	64.94	15.00
123 ARG O	-39.01	-35.82	65.89	15.00
124 ALA N	-37.15	-36.31	64.71	15.00
124 ALA CA	-36.22	-35.65	65.62	15.00
124 ALA CB	-34.86	-35.50	64.98	15.00
124 ALA C	-36.70	-34.29	66.11	15.00
124 ALA O	-36.67	-34.02	67.31	15.00
125 VAL N	-37.16	-33.45	65.19	15.00
125 VAL CA	-37.66	-32.13	65.55	15.00
125 VAL CB	-38.00	-31.28	€4.27	15.00
125 VAL CG1	-38.50	-29.89	64.64	15.00

125 VAL CG2	-36.77	-31.11	63.41	15.00
125 VAL C	-38.87		66.47	15.00
125 VAL 0	-39.03		67.41	15.00
126 ALA N	-39.71		66.28	15.00
126 ALA CA	-40.90		67.11	15.00
126 ALA CB	-42.00		66.32	15.00
126 ALA C	-40.64		68.44	15.00
126 ALA O	-41.32		69.44	15.00
127 ARG N	-39.67			15.00
127 ARG CA	-39.36		69.68	15.00
127 ARG CB	-38.79	-37.22	69.36	15.00
127 ARG CG	-39.80		69.36	15.00
127 ARG CD	-40.24	-38.71	67.96	15.00
127 ARG NE	-39.78		67.58	15.00
127 ARG CZ	-40.46	-40.88	66.80	15.00
127 ARG NH1	-39.94	-42.06	66.52	15.00
127 ARG NH2	-41.66	-40.57	66.33	15.00
127 ARG C	-38.36	-35.09	70.56	15.00
127 ARG O	-38.41	-35.20	71.78	15.00
128 VAL N	-37.44	-34.37	69.94	15.00
128 VAL CA	-36.42	-33.64	70.68	15.00
128 VAL CB	-35.07	-33.78	69.99	15.00
128 VAL CG1	-33.97	-33.24	70.88	15.00
128 VAL CG2	-34.81	-35.23	69.64	15.00
128 VAL C	-36.75	-32.17	70.84	15.00
128 VAL O	-36.91	-31.68	71.94	15.00
129 GLY N	-36.85	-31.47	69.72	15.00
129 GLY CA	-37.13	-30.05	69.75	15.00
129 GLY C	-36.29	-29.43	68.64	15.00
129 GLY 0		-30.14	67.70	15.00
130 PRO N		-28.15	68.76	15.00
130 PRO CD 130 PRO CA		-27.22	69.86	15.00
130 PRO CA		-27.49	67.75	15.00
	-34.80		68.39	15.00
130 PRO CG	-36.01		69.20	15.00
130 PRO C 130 PRO O		-28.24	67.40	15.00
131 VAL N		-28.57	68.28	15.00
131 VAL N		-28.47	66.11	15.00
131 VAL CA		-29.17	65.62	15.00
131 VAL CB		-30.36	64.73	15.00
131 VAL CG1		-31.12	64.19	
131 VAL CG2		-31.29		
TOT AWT C	-31.59	-28.20	64.81	15.00

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131	VAL	0	-32.10	-27.31	64.12	15.00
132	SER	N	-30.28	-28.33	64.94	15.00
132	SER	CA	-29.35	-27.48	64.21	15.00
132	SER	CB	-28.02	-27.41	64.95	15.00
132	SER	OG	-28.18	-26.80	66.21	15.00
132	SER	С	-29.16	-28.09	62.83	15.00
132	SER	0	-28.84	-29.27	62.71	15.00
133	VAL	N	-29.37	-27.32	61.78	15.00
133	VAL	CA	-29.21	-27.82	60.42	15.00
133	VAL	CB	-30.58	-28.02	59.71	15.00
133	VAL	CG1	-31.50	-28.91	60.54	15.00
133	VAL	CG2	-31.24	-26.68	59.44	15.00
133	VAL	C	-28.37	-26.84	59.60	15.00
133	VAL	0	-28.20	-25.68	59.99	15.00
	ALA	N	-27.82	-27.30	58.48	15.00
134		CA	-27.00	-26.45	57.62	15.00
134			-25.58	-26.95	57.57	15.00
	ALA		-27.65	-26.50	56.26	15.00
	ALA		-28.14	-27.55	55.86	15.00
135	ILE		-27.66	-25.38	55.55	15.00
135	ILE		-28.30	-25.28	54.23	15.00
135	ILE		-29.70	-24.57	54.29	15.00
135	ILE		-30.71	-25.39	55.10	15.00
135	ILE		-29.54	-23.15	54.87	15.00
135	ILE		-30.81	-22.34	54.88	15.00
135	ILE		-27.46	-24.44	53.29	15.00
135	ILE		-26.40	-23.92	53.66	15.00
	ASP		-27.97	-24.32	52.07	15.00
	ASP		-27.34	-23.50	51.04	15.00
	ASP		-27.56	-24.11	49.65	15.00
136			-27.02	-23.23	48.53	15.00
	ASP		-27.80	-22.84	47.64	15.00
	ASP		-25.83	-22.91	48.54	15.00
	ASP			-22.16	51.14	15.00
	ASP		-29.21	-22.03	50.74	15.00
	ALA		-27.43	-21.17	51.76	15.00
	ALA		-28.07	-19.87	51.86	15.00
	ALA		-27.97	-19.34	53.26	15.00
	ALA		-27.47	-18.89	50.85	15.00
	ALA		-27.92	-17.75	50.72	15.00
138	SER		-26.43	-19.33	50.15	15.00
	SER		-25.79	-18.51	49.14	15.00
138	SER	CB	-24.36	-19.00	48.90	15.00

138	SER	OG	-23.55	-18.75	50.04	15.00
138	SER	С	-26.61	-18.61	47.87	15.00
138	SER	0	-26.41	-19.53	47.07	15.00
139	LEU	N	-27.56	-17.70	47.74	15.00
139	LEU	CA	-28.46	-17.66	46.60	15.00
139	LEU	CB	-29.32	-18.92	46.55	15.00
139	LEU	CG	-30.03	-19.30	45.25	15.00
139	LEU	CD1	-29.04	-19.96	44.31	15.00
139	LEU	CD2	-31.14	-20.28	45.54	15.00
139	LEU	С	-29.35	-16.43	46.82	15.00
139	LEU	0	-29.99	-16.29	47.87	15.00
140	THR	N	-29.39	-15.54	45.83	15.00
140	THR	CA	-30.17	-14.30	45.90	15.00
140	THR	CB	-30.15	-13.57	44.52	15.00
140	THR	0G1	-29.95	-14.53	43.47	15.00
140	THR	CG2	-29.05	-12.52	44.48	15.00
140	THR	C	-31.60	-14.43	46.41	15.00
140	THR		-32.02	-13.70	47.31	15.00
141	SER	N	-32.35	-15.38	45.88	15.00
141	SER	CA	-33.74	-15.58	46.27	15.00
141	SER	CB	-34.42	-16.62	45.36	15.00
141	SER	OG	-33.62	-17.78	45.21	15.00
141	SER	С	-33.92	-15.96	47.74	15.00
141	SER	0	-34.99	-15.74	48.33	15.00
142	PHE	N	-32.90	-16.52	48.37	15.00
142	PHE	CA	-33.01	-16.91	49.77	15.00
142	PHE		-31.92	-17.91	50.15	15.00
142	PHE		-31.91	-18.26	51.61	15.00
142	PHE		-32.74		52.10	15.00
142	PHE		-31.09	-17.58	52.50	15.00
142		CE1	-32.74		53.45	15.00
142	PHE		-31.09	-17.90	53.87	15.00
	PHE		-31.92	-18.89	54.34	15.00
	PHE			-15.67	50.62	15.00
	PHE			-15.45	51.55	15.00
	GLN			-14.85	50.24	15.00
			-31.58	-13.63	50.96	15.00
	GLN		~30.25	-13.12	50.48	15.00
	GLN			-1.4.20	50.55	15.00
	GLN			-13.73	50.06	15.00
	GLN		-27.33		50.59	15.00
	GIN			14.40	49.05	15.00
143	GLN	С	-32.63	·-12.53	50.92	15.00

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143	GLN	0	-32.79	-11.82	51.91	15.00
144	PHE	N	-33.31	-12.35	49.79	15.00
144	PHE	CA	-34.36	-11.32	49.70	15.00
144	PHE	CB	-34.28	-10.50	48.39	15.00
144	PHE	CG	-34.49	-11.30	47.11	15.00
144	PHE	CD1	-33.53	-11.26	46.11	15.00
144	PHE	CD2	-35.66	-12.02	46.88	15.00
144	PHE	CE1	-33.74	-11.93	44.90	15.00
144	PHE	CE2	-35.88	-12.69	45.67	15.00
144	PHE	CZ	-34.91	-12.64	44.68	15.00
144	PHE	C	-35.77	-11.88	49.96	15.00
144	PHE	0	-36.77	-11.36	49.45	15.00
145	TYR	N	-35.82	-12.95	50.76	15.00
145	TYR	CA	-37.05	-13.64	51.13	15.00
145	TYR	CB	-36.69	-14.96	51.83	15.00
145	TYR	CG	-37.83	-15.59	52.60	15.00
145	TYR	CD1	-38.64	-16.56	52.01	15.00
145	TYR	CE1	-39.71	-17.11	52.70	15.00
145	TYR	CD2	-38.11	-15.20	53.91	15.00
145	TYR	CE2	-39.18	-15.74	54.60	15.00
145	TYR	CZ	-39.98	-16.69	53.99	15.00
145	TYR	OH	-41.05	-17.22	54.66	15.00
145	TYR	С	-37.79	-12.74	52.10	15.00
145	TYR	0	-37.16	-12.04	52.89	15.00
146	SER	N	-39.12	-12.80	52.09	15.00
146	SER	CA	-39.93	-11.97	52.99	15.00
146	SER	CB	-40.22	-10.61	52.35	15.00
146	SER	<b>OG</b>	-40.75	-10.75	51.05	15.00
146	SER	С	-41.25	-12.62	53.43	15.00
146	SER	0	-41.83	-12.21	54.43	15.00
147	LYS	N	-41.74	-13.59	52.66	15.00
147	LYS			-14.28	52.99	15.00
	LYS		-44.19	-13.35	52.82	15.00
	LYS	CG	-44.40	-12.86	51.40	15.00
147			-45.58	-11.90	51.31	15.00
147	LYS	CE	-46.86	-12.63	50.95	15.00
147	LYS	NZ	-48.03	-11.69	50.96	15.00
147	LYS	С	-43.18	-15.52	52.13	15.00
147	LYS		-42.61	-15.62	51.04	15.00
148	GLY		-44.00	-16.45	52 . 62	15.00
148	GLY	CA	-44.27	-17.68	51.88	15.00
148			-43.30	-18.31	52.16	15.00
148	GLY	0	-42.38	-18.69	52.97	15.00

149 VAL N	-43.52	-19.94	51.51	15.00
149 VAL CA	-42.66	-21.10	51.68	15.00
149 VAL CB	-43.47	-22.41	51.46	15.00
149 VAL CG1	-42.59	-23.63	51.61	15.00
149 VAL CG2	-44.62	-22.47	52.45	15.00
149 VAL C	-41.57	-20.96	50.62	15.00
149 VAL 0	-41.84	-20.56	49.50	15.00
150 TYR N	-40.34	-21.30	50.96	15.00
150 TYR CA	-39.24	-21.14	50.02	15.00
150 TYR CB	-38.02	-20.54	50.73	15.00
150 TYR CG	-36.80	-20.39	49.85	15.00
150 TYR CD1	-36.77	-19.48	48.80	15.00
150 TYR CE1	-35.66	-19.36	47.97	15.00
150 TYR CD2	-35.67	-21.18	50.05	15.00
150 TYR CE2	-34.56	-21.07	49.24	15.00
150 TYR CZ	-34.56	-20.16	48.20	15.00
150 TYR OH	-33.45	-20.04	47.40	15.00
150 TYR C	-38.83	-22.39	49.27	15.00
150 TYR O	-38.66	-23.45	49.85	15.00
151 TYR N	-38.62	-22.22	47.97	15.00
151 TYR CA	-38.17	-23.31	47.12	15.00
151 TYR CB	-39.33	-24.21	46.71	15.00
151 TYR CG	-38.86	-25.44	45.98	15.00
151 TYR CD1	-37.95	-26.30	46.56	15.00
151 TYR CE1	-37.46	-27.40	45.87	15.00
151 TYR CD2	-39.29	-25.71	44.68	15.00
151 TYR CE2	-38.81	-26.81	43.98	15.00
151 TYR CZ	-37.89	-27.65	44.58	15.00
151 TYR OH	-37.37	-28.73	43.90	15.00
151 TYR C	-37.49	-22.74	45.88	15.00
151 TYR 0	-37.97	-21.79	45.28	15.00
152 ASP N	-36.36		45.49	15.00
152 ASP CA	-35.64		44.31	15.00
152 ASP CB	-34.72	-21.69	44.66	15.00
		-21.09		
152 ASP OD1		-21.69		
152 ASP OD2		-20.01	43.01	15.00
152 ASP C	-34.83		43.66	15.00
152 ASP 0		-24.45	44.23	15.00
153 GLU N		-24.32		
153 GLU CA		-25.36		
153 GLU CB		-25.35		
153 GLU CG	-35.22	-23.98	3 <b>9.55</b>	15.00

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153 GLU CD			38.93	15.00
153 GLU OE1		3 -22.41	39.56	
153 GLU 0E2	-33.56	-23.73	37.80	
153 GLU C	-33.02	-25.27	41.80	
153 GLU 0	-32.36			
154 SER N	-32.48	-24.05	41.74	
154 SER CA	-31.02		41.81	
154 SER CB	-30.65		41.31	
154 SER OG	~30.66		39.90	
154 SER C	-30.37		43.16	15.00
154 SER 0	-29.14		43.30	
155 CYS N	-31.15		44.19	
155 CYS CA	-30.56	-24.61	45.49	
155 CYS C	-29.70	-25.86	45.37	15.00
155 CYS 0	-30.17	-26.88	44.86	15.00
155 CYS CB	-31.63	-24.79	46.55	15.00
155 CYS SG	-31.06		48.11	15.00
156 ASN N	-28.43		45.74	15.00
156 ASN CA	-27.50	-26.87	45.66	15.00
156 ASN CB	-26.13	-26.39	45.18	15.00
156 ASN CG	-25.14	-27.52	44.97	15.00
156 ASN OD1	-25.51	-28.70	44.95	15.00
156 ASN ND2	-23.88	-27.16	44.80	15.00
156 ASN C	-27.34	-27.66	46.95	15.00
156 ASN 0	-26.57	-27.31	47.85	15.00
157 SER N	-28.03	-28.79	46.98	15.00
157 SER CA	-28.03	-29.71	48.10	15.00
157 SER CB	-28.72	-31.00	47.65	15.00
157 SER OG	-29.25	-30.85	46.33	15.00
157 SER C	-26.63	-30.03	48.63	15.00
157 SER 0	-26.46	-30.47	49.77	15.00
158 ASP N	-25.61	-29.83	47.80	15.00
158 ASP CA	-24.23	-30.12	48.18	15.00
158 ASP CB	-23.50	-30.73	47.00	15.00
158 ASP CG	-23.99	-32.13	46.66	15.00
158 ASP OD1	-24.82	-32.27	45.74	15.00
158 ASP OD2	-23.57	-33.09	47.34	15.00
158 ASP C	-23.44	-28.95	48.72	15.00
158 ASP O		-29.13	49.48	15.00
159 ASN N		-27.73	48.33	15.00
159 ASN CA		-26.56		15.00
159 ASN CB	-23.07		47.75	15.00
159 ASN CG		-24 26	48.11	15.00

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# TABLE II

150		_		
159 ASN OD1				15.00
159 ASN ND2			47.22	15.00
159 ASN C	-23.63		50.13	15.00
159 ASN 0	-24.43	·	50.17	15.00
160 LEU N	-23.27		51.23	15.00
160 LEU CA	-23.78		52.55	15.00
160 LEU CB	-23.76		53.53	15.00
160 LEU CG	-24.57		53.10	15.00
160 LEU CD1	-24.08	· -	53.84	15.00
160 LEU CD2	-26.06		53.32	15.00
160 LEU C	-22.87		53.02	15.00
160 LEU O	-21.70		53.29	15.00
161 ASN N	-23.41		53.16	
161 ASN CA	-22.59		53.54	
161 ASN CB	-22.43	-21.97	52.31	15.00
161 ASN CG	-23.75	-21.64	51.67	15.00
161 ASN OD1	-24.51	-20.79	52.17	15.00
161 ASN ND2	-24.09	-22.36	50.62	15.00
161 ASN C	-23.07	-22.00	54.70	15.00
161 ASN 0	-22.32	-21.20	55.27	15.00
162 HIS N	-24.34	-22.10	55.06	15.00
162 HIS CA	-24.87	-21.31	56.16	15.00
162 HIS CB	-25.90	-20.34	55.60	15.00
162 HIS CG	-26.42	-19.36	56.60	15.00
162 HIS CD2	-27.67		56.94	15.00
162 HIS ND1	-25.58	-18.61	57.40	15.00
162 HIS CE1	-26.30	-17.82	58.18	15.00
162 HIS NE2	-27.57	-18.05	57.92	15.00
162 HIS C	-25.52	-22.24	57.17	15.00
162 HIS O	-26.26	-23.14	56.80	15.00
163 ALA N	-25.22	-22.02	58.45	15.00
163 ALA CA	-25.79	-22.84	59.53	15.00
163 ALA CB	-24.77	-23.10	60.61	15.00
163 ALA C	-27.00	-22.13	60.10	15.00
163 ALA O	-26.93		60.48	15.00
164 VAL N	-28.10	-22.85	60.20	15.00
164 VAL CA		-22.30	60.70	15.00
164 VAL CB		-22.02	59.52	15.00
164 VAL CG1		-23.26	59.13	15.00
164 VAL CG2		-20.83	59.80	15.00
164 VAL C		-23.25	61.73	15.00
164 VAL O		-24.21		15.00
165 LEU N		-22.99	62.16	15.00

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165 LEU CA	-31.87		63.17	15.00
165 LEU CB	-31.87		64.52	
165 LEU CG	-32.48	-23.78	65.74	
165 LEU CD1	-31.58	-24.90	66.23	15.00
165 LEU CD2	-32.71	-22.77		15.00
165 LEU C	-33.31	-24.16	62.81	15.00
165 LEU O	-34.08	-23.27	62.49	15.00
166 ALA N	-33.69	-25.43	62.90	15.00
166 ALA CA	-35.06	-25.86	62.59	15.00
166 ALA CB	-35.05	-27.27	62.00	15.00
166 ALA C	-35.91	-25.81	63.86	15.00
166 ALA O	-35.69	-26.57	64.80	15.00
167 VAL N	-36.89	-24.92	63.85	15.00
167 VAL CA	-37.78	-24.65	65.00	15.00
167 VAL CB	-37.89	-23.09	65.18	15.00
167 VAL CG1	-38.97	-22.71	66.15	15.00
167 VAL CG2	-36.58	-22.55	65.68	15.00
167 VAL C	-39.16	-25.32	64.93	15.00
167 VAL O	-39.98	-25.21	65.84	15.00
168 GLY N	-39.43	-26.06	63.87	15.00
168 GLY CA	-40.71	-26.72	63.75	15.00
168 GLY C	-40.98	-27.09	62.31	15.00
168 GLY O	-40.05	-27.10	61.49	15.00
169 TYR N	-42.23	-27.39	61.99	15.00
169 TYR CA	-42.65	-27.76	60.65	15.00
169 TYR CB	-42.15	-29.17	60.29	15.00
169 TYR CG	-42.64	-30.27	61.22	15.00
169 TYR CD1	-44.00	-30.58	61.31	15.00
169 TYR CE1	-44.46	-31.57	62.15	15.00
169 TYR CD2	-41.75	-30.99	62.01	15.00
169 TYR CE2	-42.20	-31.99	62.85	15.00
169 TYR CZ	-43.56	-32.28	62.91	15.00
169 TYR OH	-44.04	-33.28	63.71	15.00
169 TYR C	-44.16	-27.70	60.54	15.00
169 TYR O	-44.85	-27.38	61.52	15.00
170 GLY N	-44.70	~28.04	59.38	15.00
170 GLY CA	-46.13	-28.00	59.18	15.00
170 GLY C		-27.76	57.73	15.00
170 GLY O		-28.26	56.83	15.00
171 ILE N		-25.92	57.48	15.00
171 ILE CA		-26.63	56.13	15.00
171 ILE CB		-27.66	55.72	15.00
171 ILE CG2		-27.09	54.71	15.00
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171 ILE CG1			55.21	15.00
171 ILE CD1			54.82	
171 ILE C	-48.54		56.14	
171 ILE 0	-48.91		57.20	
172 GLN N	-48.58			
172 GLN CA	-49.16			
172 GLN CB	-48.08	-22.18	54.62	15.00
172 GLN CG	-48.58	_		
172 GLN CD	-47.50	_		
172 GLN OE1	-46.95			
172 GLN NE2	-47.18	-18.82	54.96	15.00
172 GLN C	-50.29	-23.15	53.96	15.00
172 GLN 0	-51.45	-22.89	54.32	15.00
173 LYS N	-49.96	-23.36	52.68	
173 LYS CA	-50.96		51.61	15.00
173 LYS CB	-50.84	-22.14	50.69	15.00
173 LYS CG	-51.09		51.36	15.00
173 LYS CD	-52.40	-20.72	52.16	15.00
173 LYS CE	-53.63	-20.91	51.29	15.00
173 LYS NZ	-54.88	-20.56	52.06	15.00
173 LYS C	-50.66	-24.63	50.81	15.00
173 LYS O		-24.57	49.67	15.00
174 GLY N	-50.86	-25.77	51.45	15.00
174 GLY CA	-50.60	-27.04	50.79	15.00
174 GLY C	-49.13	-27.39	50.91	15.00
174 GLY O	-48.77	-28.52	51.28	15.00
175 ASN N	-48.26	-26.42	50.63	15.00
175 ASN CA	-46.83	-26.66	50.70	15.00
175 ASN CB	-46.06	-25.48	50.10	15.00
175 ASN CG	-46.31	-25.33	48.61	15.00
175 ASN OD1	-47.34	-24.79	48.20	15.00
175 ASN ND2	-45.38	-25.81	47.30	15.00
175 ASN C		-26.96	52.11	15.00
175 ASN 0	-46.55	-26.18	53.05	15.00
176 LYS N	-45.69	-28.13	52.25	15.00
176 LYS CA	-45.13	-28.55	53.53	15.00
176 LYS CB		-30.01	53.48	15.00
176 LYS CG	-45.77	-31.03	53.17	15.00
176 LYS CD		-32.45	53.44	15.00
176 LYS CE	-46.19	-33.50	52.85	15.00
176 LYS NZ		-33.64	51.36	15.00
176 LYS C	-43.92	-27.65	53.74	15.00
176 LYS O		-27.27	52.77	15.00

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177	HIS	N	-43.57	-27.37	54.99	15.00
177	HIS	CA	-42.44	-26.49	55.22	15.00
177	HIS	CB	-42.90	-25.04	55.10	15.00
177	HIS	CG	-43.81	-24.59	56.20	15.00
177	HIS	CD2	-43.59	-24.40	57.52	15.00
177	HIS	ND1	-45.13	-24.24	55.98	15.00
177	HIS	CE1	-45.67	-23.85	57.12	15.00
177	HIS	NE2	-44.76	-23.93	58.07	15.00
177	HIS	C	-41.73	-26.66	56.55	15.00
177	HIS	0	-42.31	-27.17	57.50	15.00
178	TRP	N	-40.48	-26.20	56.61	15.00
178	TRP	CA	-39.68	-26.25	57.82	15.00
178	TRP	CB	-38.26	-26.73	57.52	15.00
178	TRP	CG	-38.13	-28.11	57.02	15.00
178	TRP	CD2	-38.29	-29.32	57.78	15.00
178	TRP	CE2	-37.96	-30.39	56.93	15.00
178	TRP	CE3	-38.68	-29.59	59.10	15.00
178	TRP	CD1	-37.74	-28.49	55.78	15.00
178	TRP	NE1	-37.62	-29.86	55.72	15.00
178	TRP	CZ2	-38:00	-31.71	57.35	15.00
178	TRP	CZ3	-38.72	-30.91	59.52	15.00
178	TRP	CH2	-38.38	-31.96	58.64	15.00
178	TRP	C	-39.60	-24.81	58.34	15.00
178	TRP	0	-39.21	-23.90	57.60	15.00
179	ILE	N	-39.96	-24.57	59.59	15.00
179	ILE	CA	-39.89	-23.23	60.17	15.00
179	ILE	CB	-40.79	-23.11	61.41	15.00
179		CG2	-40.66	-21.74	62.03	15.00
179	ILE	CG1	-42.24	-23.42	61.02	15.00
179	ILE	CD1	-43.21	-23.40	62.15	15.00
179		С	-38.44	-23.00	60.58	15.00
179	ILE		-37.97	-23.57	61.56	15.00
180	ILE		-37.72	-22.17	59.81	15.00
	ILE	CA	-36.31	-21.89	60.07	15.00
180	ILE		-35.49	-21.93	58.76	15.00
180	ILE		-34.04	-21.77	59.05	15.00
	ILE		-35.73	-23.25	58.02	15.00
	ILE		-35.30	-24.46	58.78	15.00
	II.E		-36.03	-20.55	60.77	15.00
180	ILE			-19.52	60.42	15.00
				-20.57	61.74	15.00
	LYS			-19.40	62.50	15.00
181	LYS	CB	-34.63	-19.75	63.99	15.00

181 LYS CG	-34.15		64.87	15.00
181 LYS CD	-33.79		66.29	15.00
181 LYS CE	-33.65	-17.84	67.19	15.00
181 LYS NZ	-33.21	-18.18	68.57	15.00
181 LYS C	-33.36	-19.00	62.01	15.00
181 LYS O	-32.43	-19.79	62.05	15.00
182 ASN N	-33.20	-17.77	61.53	15.00
182 ASN CA	-31.90	-17.34	61.04	15.00
182 ASN CB	-32.07	-16.64	59.69	15.00
182 ASN CG	-30.84	-16.74	58.82	15.00
182 ASN OD1	-29.75	-17.06	59.30	15.00
182 ASN ND2	-31.01	-16.50	57.53	15.00
182 ASN C	-31.22	-16.43	62.06	15.00
182 ASN 0	-31.77	-16.14	63.11	15.00
183 SER N	-30.00	-15.98	61.77	15.00
183 SER CA	-29.29	-15.10	62.69	15.00
183 SER CB	-28.07	-15.81	63.25	15.00
183 SER OG	-27.40	-16.52	62.23	15.00
183 SER C	-28.87	-13.82	61.98	15.00
183 SER O	-27.82	-13.25	62.27	15.00
184 TRP N	-29.70	-13.35	61.06	15.00
184 TRP CA	-29.40	-12.13	60.34	15.00
184 TRP CB	-29.58	-12.36	58.83	15.00
184 TRP CG	-28.57	-13.29	58.22	15.00
184 TRP CD2	-28.61	-13.87	56.90	15.00
184 TRP CE2	-27.40	-14.58	56.72	15.00
184 TRP CE3	-29.54	-13.84	55.86°	15.00
184 TRP CD1	-27.39	-13.69	58.76	15.00
184 TRP NE1	-26.68	-14.46	57.87	15.00
184 TRP CZ2	-27.11	-15.26	55.53	15.00
184 TRP CZ3	-29.25	-14.52	54.67	15.00
184 TRP CH2	-28.04	-15.22	54.52	15.00
184 TRP C	-30.28	-10.98	60.82	15.00
184 TRP O	-30.61	-10.09	60.04	15.00
185 GLY N	-30.70	-11.02	62.09	15.00
185 GLY CA	-31.53	-9.96	62.66	15.00
L85 GLY C	-33.03	-10.14	62.46	15.00
L85 GLY O	-33.46	-10.99	61.69	15.00
186 GLU N	-33.84	-9.34	63.17	15.00
L86 GLU CA	-35.30	-9.44	63.04	15.00
186 GLU CB	-36.00	-8.71	64.19	1500
186 GLU CG	-35.52	-9.13	65.56	15.00
.86 GLU CD	-36.52	-8.85	66.68	15.00

186 GLU	OE1	-36.30	-9.38	67.80	15.00
186 GLU	OE2	-37.51	-8.13	66.46	15.00
186 GLU	C	-35.74	-8.83	61.73	15.00
186 GLU	0	-36.83	-9.11	61.22	15.00
187 ASN	N	-34.89	-7.97	61.19	15.00
187 ASN	CA	-35.15	-7.28	59.95	15.00
187 ASN	CB	-34.04	-6.25	59.73	15.00
187 ASN	CG	-34.56	-4.94	59.17	15.00
187 ASN	OD1	-33.77	-4.07	58.80	15.00
187 ASN	ND2	-35.88	-4.76	59.16	15.00
187 ASN	С	-35.21	-8.22	58.75	15.00
187 ASN	0	-36.04	-8.04	57.86	15.00
188 TRP	N	-34.33	-9.22	58.73	15.00
188 TRP	CA	-34.26	-10.18	57.64	15.00
188 TRP	CB	-33.03	-11.07	57.79	15.00
188 TRP	CG	-32.85	-12.02	56.65	15.00
188 TRP	CD2	-33.28	-13.38	56.57	15.00
188 TRP	CE2	-32.98	-13.85	55.28	15.00
188 TRP	CE3	-33.90	-14.26	57.48	15.00
188 TRP	CD1	-32.31	-11.72	55.44	15.00
188 TRP	NE1	-32.39	-12.81	54.61	15.00
188 TRP		-33.27	-15.15	54.86	15.00
188 TRP		-34.20	-15.55	57.06	15.00
188 TRP	CH2	-33.88	-15.98	55.77	15.00
188 TRP	С	-35.50	-11.05	57.53	15.00
188 TRP		-36.10	-11.42	58.55	15.00
189 GLY 1		-35.85	-11.42	56.31	15.00
189 GLY (	CA	-37.00	-12.26	56.07	15.00
189 GLY (		-38.21	-11.92	56.90	15.00
189 GLY (		-38.47	-10.76	57.23	15.00
190 ASN 1		-38.97	-12.94	57.27	15.00
190 ASN (		-40.16	-12.73	58.07	15.00
190 ASN (		-41.17	-13.86	57.83	15.00
190 ASN (		-42.55		58.36	15.00
190 ASN (		-42.70	-12.88	59.39	15.00
190 ASN 1		-43.57	-13.99	57.65	15.00
190 ASN (		-39.79	-12.67	59.54	15.00
190 ASN (		-39.85	-13.66	60.25	15.00
191 LYS 1		-39.32	-11.52	60.00	15.00
191 LYS (		-38.95	-11.34	61.39	15.00
191 LYS (				62.29	15.00
191 LYS (	CG	-40.95	-10.18	62.44	15.00
191 LYS (	CD	-42.19	-10.32	63.30	15.00

101			_	
191 LYS CE	-43.32			15.00
191 LYS NZ	-43.71			15.00
191 LYS C	-37.84			15.00
191 LYS O	-37.77			15.00
192 GLY N	-36.94			15.00
192 GLY CA	-35.83		61.35	15.00
192 GLY C	-36.05		60.98	15.00
192 GLY O	-35.12		60.99	15.00
193 TYR N	-37.29		60.65	15.00
193 TYR CA	-37.60		60.28	15.00
193 TYR CB	-38.84		61.05	15.00
193 TYR CG	-38.60		62.52	15.00
193 TYR CD1	-38.73		63.41	15.00
193 TYR CE1	-38.46		64.76	15.00
193 TYR CD2	-38.19		63.03	15.00
193 TYR CE2	-37.92		64.38	15.00
193 TYR CZ	-38.05		65.23	15.00
193 TYR OH	-37.77	-17.80	66.56	15.00
193 TYR C	-37.83	-16.81	58.79	15.00
193 TYR O	-38.02	-15.82	58.08	15.00
194 ILE N	-37.82	-18.06	58.35	15.00
194 ILE CA	-38.04	-18.41	56.95	15.00
194 ILE CB	-36.75	-18.32	56.10	15.00
194 ILE CG2	-35.64	-19.14	56.72	15.00
194 ILE CG1	-37.03	-18.83	54.69	15.00
194 ILE CD1	-35.86	-18.72	53.75	15.00
194 ILE C	-38.61	-19.81	56.83	15.00
194 ILE 0	-38.11	-20.74	57.43	15.00
195 LEU N	-39.72	-19.92	56.10	15.00
195 LEU CA	-40.38	-21.19	55.85	15.00
195 LEU CB	-41.88	-20.96	55.65	15.00
195 LEU CG	-42.82	-20.84	56.86	15.00
195 LEU CD1	-42.19	-20.08	58.01	15.00
195 LEU CD2	-44.12	-20.18	56.42	15.00
195 LEU C	-39.76	-21.78	54.59	15.00
195 LEU O	-39.79	-21.16	53.53	15.00
196 MET N	-39.11	-22.93	54.71	15.00
196 MET CA	-38.46	-23.57	53.57	15.00
196 MET CB	-37.03	-23.97	53.96	15.00
196 MET CG		-22.77	54.14	15.00
196 MET SD		-23.18	54.74	15.00
196 MET CE		-23.82	53.22	15.00
196 MET C		-24.78	53.11	15.00
		-		

	MET		-40.09	-25.31	53.86	15.00
197			-39.08	-25.23	51.87	15.00
197	ALA	CA	-39.83	-26.37	51.32	15.00
197	ALA	CB	-39.62	-26.47	49.82	15.00
197	ALA	С	-39.52	-27.71	51.98	15.00
197	ALA	0	-38.37	-28.15	51.98	15.00
198	ARG	N	-40.56	-28.39	52.46	15.00
198	ARG	CA	-40.40	-29.68	53.11	15.00
198	ARG	CB	-41.17	-29.72	54.44	15.00
198	ARG	CG	-41.15	-31.07	55.15	15.00
198	ARG	CD	-41.43	-30.95	56.63	15.00
198	ARG	NE	-42.81	-30.59	56.92	15.00
198	ARG	CZ	-43.79	-31.47	57.10	15.00
198	ARG	NH1	-43.54	-32.78	57.01	15.00
198	ARG	NH2	-45.01	-31.05	57.42	15.00
198	ARG	С	-40.82	-30.85	52.23	15.00
198	ARG	0	-41.86	-30.80	51.57	15.00
199	asn		-40.00	-31.89	52.23	15.00
199	ASN		-40.25	-33.10	51.45	15.00
199	ASN		-41.59	-33.75	51.83	15.00
199	ASN	CG	-41.57	-34.40	53.22	15.00
199	ASN	OD1	-42.60	-34.43	53.91	15.00
199	ASN	ND2	-40.42	-34.91	53.63	15.00
199	ASN	С	-40.15	-32.94	49.94	15.00
199	asn	0	-40.49	-33.86	49.19	15.00
200	LYS	N	-39.67	-31.77	49.49	15.00
200	LYS	CA	-39.50	-31.55	48.05	15.00
200	LYS	CB	-39.77	-30.09	47.65	15.00
200	LYS	CG	-41.23	-29.79	47.39	15.00
200		CD	-41.42	-28.49	46.63	15.00
200	LYS	CE	-42.88	-28.03	46.65	15.00
200		NZ	-43.06	-26.68	46.03	15.00
200	LYS	C	-38.09	-31.95	47.69	15.00
	LYS		-37.28	-31.12	47.26	15.00
	ASN		-37.80	-33.22	47.95	15.00
201	ASN	CA	-36.50	-33.81	47.65	15.00
201	ASN	CB	-36.24	-33.83	46.13	15.00
	ASN	CG	-37.32	-34.57	45.35	15.00
		OD1	-37.09	-35.68	44.87	15.00
	ASN		-38.46	-3392	45.15	15.00
201	ASN	С	-35.31	-33.16	48.37	15.00
201	ASN	0	-34.31	-32.84	47.74	15.00
202	ASN	N	-35.43	-32.90	49.67	15.00

202 ASN CA	-34.32	-32.33	50.44	15.00
202 ASN CB	-33.19	-33.35	50.50	15.00
202 ASN CG	-32.27	-33.14	51.68	15.00
202 ASN OD1	-32.71	-32.70	52.75	15.00
202 ASN ND2	-31.00	-33.48	51.51	15.00
202 ASN C	-33.81	-30.99	49.89	15.00
202 ASN O	-32.60		49.75	15.00
203 ALA N	-34.73	-30.09	49.59	15.00
203 ALA H	-35.63	-30.29	49.94	15.00
203 ALA CA	-34.40	-28.79	49.02	15.00
203 ALA CB	-35.62	-27.88	48.98	15.00
203 ALA C	-33.35	-28.09	49.90	15.00
203 ALA O	-33.51	-27.93	51.10	15.00
204 CYS N	-32.27	-27.66	49.25	15.00
204 CYS CA	-31.18	-26.94	49.91	15.00
204 CYS C	-30.38	-27.79	50.89	15.00
204 CYS O	-29.60	-27.25	51.68	15.00
204 CYS CB	-31.71	-25.68	50.59	15.00
204 CYS SG	-32.51	-24.48	49.47	15.00
205 GLY N	-30.56	-29.10	50.84	15.00
205 GLY CA	-29.83	-30.00	51.71	15.00
205 GLY C	-30.11	-29.81	53.19	15.00
205 GLY O	-29.22	-29.98	54.02	15.00
206 ILE N	-31.35	-29.51	53.52	15.00
206 ILE CA	-31.77	-29.30	54.89	15.00
206 ILE CB	-33.30	-29.10	54.96	15.00
206 ILE CG2	-34.02	-30.31	54.38	15.00
206 ILE CG1	-33.76	-28.86	56.41	15.00
206 ILE CD1	-33.48	-27.47	56.92	15.00
206 ILE C	-31.35	-30.45	55.82	15.00
206 ILE O	-30.94	-30.23	56.97	15.00
207 ALA N	-31.36	-31.68	55.31	15.00
207 ALA CA	-31.00	-32.84	56.11	15.00
207 ALA CB	-32.05	-33.93	55.95	15.00
207. ALA C	-29.61	-33.42	55.88	15.00
207 ALA O	-29.35	-34.57	56.24	15.00
208 ASN N	-28.68	-32.65	55.31	15.00
208 ASN CA		-33.15	55.05	15.00
208 ASN CB	-26.76	-32.53	53.77	15.00
208 ASN CG		-33.15		
208 ASN OD1	-28.21	-34.02	52.57	15.00
208 ASN ND2	-26.89	-32.67	51.36	15.00
208 ASN C	-26.32	-32.95	56.20	15.00

208 ASN 0	-25.17	-33.40	56.12	15.00
209 LEU N	-26.72	-32.25	57.25	15.00
209 LEU CA	-25.84	-32.03	58.40	15.00
209 LEU CB	-24.76	-31.00	58.04	15.00
209 LEU CG	-23.41	-30.99	58.78	15.00
209 LEU CD1	-22.72	-32.35	58.64	15.00
209 LEU CD2	-22.51	-29.86	58.24	15.00
209 LEU C	-26.67	-31.60	59.63	15.00
209 LEU O	-26.38	-30.59	60.28	15.00
210 ALA N	-27.72	-32.37	59.95	15.00
210 ALA H	-28.03	-32.90	59.19	15.00
210 ALA CA	-28.61	-32.05	61.06	15.00
210 ALA CB	-30.02	-32.57	60.81	15.00
210 ALA C	-28.10	-32.73	62.34	15.00
210 ALA O	-27.62	-33.86	62.34	15.00
211 SER N	-28.18	-32.02	63.47	15.00
211 SER CA	-27.75	-32.59	64.74	15.00
211 SER CB	-26.24	-32.46	64.92	15.00
211 SER OG	-25.84	-31.12	65.06	15.00
211 SER C	-28.45	-31.88	65.88	15.00
211 SER 0	-29.03	-30.80	65.71	15.00
212 PHE N	-28.50	-32.51	67.04	15.00
212 PHE CA	-29.11	-31.91	68.21	15.00
212 PHE CB	-30.59	-32.31	68.33	15.00
212 PHE CG	-30.81	-33.79	68.43	15.00
212 PHE CD1	-31.11	-34.54	67.30	15.00
212 PHE CD2	-30.72	-34.44	69.66	15.00
212 PHE CE1	-31.32	-35.91	67.39	15.00
212 PHE CE2	-30.92	-35.81	69.77	15.00
212 PHE CZ	-31.23	-36.55	68.63	15.00
212 PHE C	-28.30	-32.39	69.40	15.00
212 PHE 0	-27.66	-33.43	69.32	15.00
213 PRO N	-28.24	-31.58	70.46	15.00
213 PRO CD	-28.77	-30.21	70.54	15.00
213 PRO CA	-27.48	-31.92	71.67	15.00
213 PRO CB	-27.21	-30.55	72.28	15.00
213 PRO CG	-28.47	-29.82	71.97	15.00
213 PRO C	-28.27	-32.81	72.64	15.00
213 PRO 0		-32.76	72.69	15.00
214 LYS N		-33.62	73.42	15.00
214 LYS CA	-28.23		74.37	15.00
214 LYS CB	-27.67		74.28	15.00
214 LYS CG	-28.06	-36.64	73.02	15.00

214 LYS CE				_	
214 LYS NZ	214 LYS CD			73.07	15.00
214 LYS C				73.05	15.00
214 LYS O		-25.79	-39.70	72.73	15.00
215 MET N		-28.03	-33.95	75.77	15.00
215 MET CA			-	76.18	15.00
215 MET CB				76.47	15.00
215 MET CB				77.83	15.00
215 MET CG		-29.71	-31.80	77.93	
215 MET SD	215 MET CG	-28.74	-30.64	77.75	
215 MET CE	215 MET SD	-29.45	-29.00	78.06	
215 MET C	215 MET CE	-30.38	-29.32	79.56	
215 MET OT1		-29.72	-34.16	78.81	
215 MET OT2		-30.39	-35.12		
216 HOH OH2		-29.55	-33.97	80.04	
217 HOH OH2	216 НОН ОН2	-28.46	-18.77	85.58	
218 HOH OH2		-24.63	-33.99	81.97	
219 HOH OH2	218 нон он2	-31.11	-15.95	65.82	
220 HOH OH2	219 нон он2	-30.23	-19.59		
222 HOH OH2	220 нон он2	-8.58	-7.31	62.36	
222 HOH OH2	221 нон он2	-6.71	-10.79	69.96	15.00
223 HOH OH2	222 нон он2	-34.27	-22.79	70.48	
224 HOH OH2   -15.68   -8.93   63.11   15.00   225 HOH OH2   -24.93   -30.84   62.42   15.00   226 HOH OH2   -7.02   -8.27   72.29   15.00   227 HOH OH2   -13.39   -20.80   66.92   15.00   228 HOH OH2   -44.55   -30.12   50.27   15.00   229 HOH OH2   -44.14   -35.34   56.06   15.00   230 HOH OH2   -37.95   -16.02   68.44   15.00   231 HOH OH2   -36.41   -36.82   52.05   15.00   232 HOH OH2   -20.00   -36.75   62.15   15.00   233 HOH OH2   -30.13   -19.30   67.02   15.00   234 HOH OH2   -28.16   -19.22   62.41   15.00   235 HOH OH2   -22.03   -29.95   62.71   15.00   236 HOH OH2   -25.92   -8.85   75.85   15.00   237 HOH OH2   -41.00   -28.78   81.30   15.00   238 HOH OH2   -32.73   -23.15   83.59   15.00   239 HOH OH2   -35.40   -24.36   49.90   15.00   240 HOH OH2   -35.40   -24.36   49.00   15.00   241 HOH OH2   -48.40   -32.54   58.07   15.00   243 HOH OH2   -27.39   -6.75   59.53   15.00   244 HOH OH2   -27.39   -6.75   59.53   15.00   245 HOH OH2   -41.50   -14.46   65.52   15.00	223 нон он2	-16.88	-33.68		
226 HOH OH2	224 НОН ОН2	-15.68	-8.93	63.11	
226 HOH OH2	225 нон он2	-24.93	-30.84	62.42	15.00
228 HOH OH2   -44.55   -30.12   50.27   15.00   229 HOH OH2   -44.14   -35.34   56.06   15.00   230 HOH OH2   -37.95   -16.02   68.44   15.00   231 HOH OH2   -36.41   -36.82   52.05   15.00   232 HOH OH2   -20.00   -36.75   62.15   15.00   233 HOH OH2   -30.13   -19.30   67.02   15.00   234 HOH OH2   -28.16   -19.22   62.41   15.00   235 HOH OH2   -22.03   -29.95   62.71   15.00   236 HOH OH2   -25.92   -8.85   75.85   15.00   237 HOH OH2   -41.00   -28.78   81.30   15.00   238 HOH OH2   -32.73   -23.15   83.59   15.00   239 HOH OH2   -40.55   -13.35   49.90   15.00   240 HOH OH2   -35.40   -24.36   49.00   15.00   241 HOH OH2   -48.40   -32.54   58.07   15.00   242 HOH OH2   -27.39   -6.75   59.53   15.00   243 HOH OH2   -41.50   -14.46   65.52   15.00		-7.02	-8.27	72.29	
229 HOH OH2   -44.14   -35.34   56.06   15.00   230 HOH OH2   -37.95   -16.02   68.44   15.00   231 HOH OH2   -36.41   -36.82   52.05   15.00   232 HOH OH2   -20.00   -36.75   62.15   15.00   233 HOH OH2   -20.13   -19.30   67.02   15.00   234 HOH OH2   -28.16   -19.22   62.41   15.00   235 HOH OH2   -22.03   -29.95   62.71   15.00   236 HOH OH2   -25.92   -8.85   75.85   15.00   237 HOH OH2   -41.00   -28.78   81.30   15.00   238 HOH OH2   -32.73   -23.15   83.59   15.00   239 HOH OH2   -40.55   -13.35   49.90   15.00   240 HOH OH2   -35.40   -24.36   49.00   15.00   241 HOH OH2   -48.40   -32.54   58.07   15.00   242 HOH OH2   -27.39   -6.75   59.53   15.00   243 HOH OH2   -27.39   -6.75   59.53   15.00   243 HOH OH2   -41.50   -14.46   65.52   15.00	227 НОН ОН2	-13.39	-20.80	66.92	15.00
230 HOH OH2   -37.95   -16.02   68.44   15.00   231 HOH OH2   -36.41   -36.82   52.05   15.00   232 HOH OH2   -20.00   -36.75   62.15   15.00   233 HOH OH2   -30.13   -19.30   67.02   15.00   234 HOH OH2   -28.16   -19.22   62.41   15.00   235 HOH OH2   -22.03   -29.95   62.71   15.00   236 HOH OH2   -25.92   -8.85   75.85   15.00   237 HOH OH2   -41.00   -28.78   81.30   15.00   238 HOH OH2   -32.73   -23.15   83.59   15.00   239 HOH OH2   -40.55   -13.35   49.90   15.00   240 HOH OH2   -35.40   -24.36   49.00   15.00   241 HOH OH2   -48.40   -32.54   58.07   15.00   242 HOH OH2   -27.39   -6.75   59.53   15.00   243 HOH OH2   -41.50   -14.46   65.52   15.00		-44.55	-30.12	50.27	15.00
231 HOH OH2   -36.41   -36.82   52.05   15.00   232 HOH OH2   -20.00   -36.75   62.15   15.00   233 HOH OH2   -30.13   -19.30   67.02   15.00   234 HOH OH2   -28.16   -19.22   62.41   15.00   235 HOH OH2   -22.03   -29.95   62.71   15.00   236 HOH OH2   -25.92   -8.85   75.85   15.00   237 HOH OH2   -41.00   -28.78   81.30   15.00   238 HOH OH2   -32.73   -23.15   83.59   15.00   239 HOH OH2   -40.55   -13.35   49.90   15.00   240 HOH OH2   -35.40   -24.36   49.00   15.00   241 HOH OH2   -48.40   -32.54   58.07   15.00   242 HOH OH2   -27.39   -6.75   59.53   15.00   243 HOH OH2   -41.50   -14.46   65.52   15.00		-44.14	-35.34	56.06	15.00
232 HOH OH2		-37.95	-16.02	68.44	15.00
233 HOH OH2 -30.13 -19.30 67.02 15.00 234 HOH OH2 -28.16 -19.22 62.41 15.00 235 HOH OH2 -22.03 -29.95 62.71 15.00 236 HOH OH2 -25.92 -8.85 75.85 15.00 237 HOH OH2 -41.00 -28.78 81.30 15.00 238 HOH OH2 -32.73 -23.15 83.59 15.00 239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00		-36.41	-36.82	52.05	15.00
234 HOH OH2 -28.16 -19.22 62.41 15.00 235 HOH OH2 -22.03 -29.95 62.71 15.00 236 HOH OH2 -25.92 -8.85 75.85 15.00 237 HOH OH2 -41.00 -28.78 81.30 15.00 238 HOH OH2 -32.73 -23.15 83.59 15.00 239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00	. –	-20.00	-36.75	62.15	15.00
235 HOH OH2 -22.03 -29.95 62.71 15.00 236 HOH OH2 -25.92 -8.85 75.85 15.00 237 HOH OH2 -41.00 -28.78 81.30 15.00 238 HOH OH2 -32.73 -23.15 83.59 15.00 239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00			-19.30	67.02	15.00
236 HOH OH2 -25.92 -8.85 75.85 15.00 237 HOH OH2 -41.00 -28.78 81.30 15.00 238 HOH OH2 -32.73 -23.15 83.59 15.00 239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00				62.41	15.00
237 HOH OH2 -41.00 -28.78 81.30 15.00 238 HOH OH2 -32.73 -23.15 83.59 15.00 239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00		-22.03	-29.95	62.71	15.00
238 HOH OH2 -32.73 -23.15 83.59 15.00 239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00			-8.85	75.85	15.00
239 HOH OH2 -40.55 -13.35 49.90 15.00 240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00		-41.00	-28.78	81.30	15.00
240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00		-32.73	-23.15	83.59	
240 HOH OH2 -35.40 -24.36 49.00 15.00 241 HOH OH2 -48.40 -32.54 58.07 15.00 242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00	<del>_</del>		-13.35	49.90	
242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00		-35.40	-24.36	49.00	15.00
242 HOH OH2 -27.39 -6.75 59.53 15.00 243 HOH OH2 -41.50 -14.46 65.52 15.00		-48.40	-32.54	58.07	15.00
243 HOH OH2 -41.50 -14.46 65.52 15.00		-27.39	-6.75	59.53	
244 77077 2000		-41.50	-14.46	65.52	
	244 HOH OH2	-22.40	-5.47	61.33	15.00

245 HC	он он2	-33.17	-27.91	70.80	15.00
246 HC	OH OH2	-45.87	-26.25	75.72	15.00
247 HC	он онг	-12.64	-13.96	81.39	15.00
248 HC	он онг	-3.78	-18.92	74.98	15.00
249 HC	он онг	-8.03	-17.70	78.42	15.00
250 HC	H OH2	-27.41	-34.98	59.22	15.00
251 HC	н он2	-34.88	-10.94	53.71	15.00
252 HC	н он2	-32.92	-27.68	46.17	15.00
253 HC	н он2	-39.35	-16.01	44.28	15.00
254 HC	н он2	-41.38	-34.64	56.30	15.00
255 HC	н он2	-44.42	-18.35	73.08	15.00
256 HC	н он2	-32.35	-13.73	61.23	15.00
257 HC	H OH2	-39.40	-8.90	59.13	15.00
258 HC	н он2	-28.41	-8.93	68.65	15.00
259 HO	н он2	-31.58	-6.53	63.69	15.00
260 HO	н он2	-19.27	-8.48	63.41	15.00
261 HO	н он2	-33.33	-20.29	70.52	15.00
262 HO	н он2	-13.49	-22.80	78.17	15.00
263 HO	н он2	-8.72	-18.49	72.60	15.00
264 HO	н он2	-10.39	-28.70	76.32	15.00
265 HO	н он2	-20.24	-31.77	61.63	15.00
266 HO	н он2	-24.78	-46.10	72.19	15.00
267 HO	н он2	-13.26	-33.12	68.94	15.00
268 HO	H OH2	-12.60	-26.87	72.01	15.00
269 HO	н он2	-17.76	-34.32	80.14	15.00
270 HO	н он2	-22.51	-37.80	70.83	15.00
271 HO	н он2	-7.33	-12.89	66.95	15.00
272 HO	н он2	-9.75	-17.21	68.77	15.00
273 HO	н он2	-30.86	-20.40	48.59	15.00
274 HO	н он2	-25.79	-24.78	42.10	15.00
275 HO	н он2	-33.50	-37.21	50.03	15.00
276 HO	н он2	-23.21	-24.90	43.38	15.00
	H OH2	-37.83	-31.49	44.10	15.00
278 HC	H OH2	-37.02	-30.78	51.01	15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors  $(A^2)$  for the cathepsin K complex with inhibitor bis-(cbz-leucinyl)-1,3-diamino-propan-2-one.

Residue	Atom	x	Y	Z	В
1 ALA	СВ	-54.29	-33.17	65.94	15.00
1 ALA	С	-53.88	-32.69	63.50	15.00
1 ALA	0	-53.42	-33.61	62.80	15.00
1 ALA		-55.60	-34.28	64.17	15.00
1 ALA	CA	-54.93	-33.01	64.57	15.00
2 PRO		-53.52	-31.40	63.32	15.00
2 PRO		-53.99	-30.23	64.09	15.00
2 PRO		-52.52	-30.98	62.32	15.00
2 PRO		-52.49	-29.46	62.49	15.00
2 PRO		-52.83	-29.26	63.94	15.00
2 PRO		-51.13	-31.59	62.52	15.00
2 PRO		-50.62	-31.64	63.64	15.00
3 ASP 1		-50.53	-32.08	61.44	15.00
3 ASP		-49.19	-32.65	61.51	15.00
3 ASP		-48.89	-33.49	60.27	15.00
3 ASP		-49.53	-34.88	60.32	15.00
3 ASP		-49.43	-35.55	61.39	15.00
3 ASP		-50.12	-35.29	59.28	15.00
3 ASP (		-48.24	-31.46	61.55	15.00
3 ASP (		-47.60	-31.14	60.54	15.00
4 SER I		-48.16	-30.78	62.68	15.00
4 SER (		-47.29	-29.62	62.80	15.00
4 SER (		-47.99	-28.35	62.27	15.00
4 SER (		-48.14	-28.37	60.86	15.00
4 SER (		-46.84	-29.35	64.23	15.00
4 SER (		-47.54	-28.71	65.19	15.00
5 VAL 1		-45.68	-28.72	64.36	15.00
5 VAL (		-45.14	-28.35	65.65	15.00
5 VAL (		-44.25	-29.47	66.25	15.00
5 VAL (		-43.09	-29.81	65.33	15.00
5 VAL 0		-43.75	-29.04	67.62	15.00
5 VAL (		-44.36	-27.05	65.44	15.00
5 VAL C		-43.60	-26.93	64.48	15.00
6 ASP N		-44.59	-25.08	66.30	15.00
6 ASP (	CA	-43.94	-24.79	66.20	15.00

6	ASP	CB	-44.95	-23.76	65.68	15.00
6	ASP	CG	-44.35	-22.38	65.47	15.00
6	ASP	OD1	-43.14	-22.26	65.21	15.00
6	ASP	OD2	-45.11	-21.39	65.56	15.00
6	ASP	С	-43.47	-24.45	67.60	15.00
6	ASP	0	-44.25	-24.02	68.45	15.00
7	TYR	N	-42.18	-24.61	67.86	15.00
7	TYR	CA	-41.62	-24.34	69.18	15.00
7	TYR	CB	-40.24	-24.98	69.29	15.00
7	TYR	CG	-40.34	-26.48	69.38	15.00
7	TYR	CD1	-40.73	-27.10	70.57	15.00
7	TYR	CE1	-40.88	-28.46	70.65	15.00
7	TYR	CD2	-40.08	-27.28	68.27	15.00
7	TYR	CE2	-40.23	-28.65	68.34	15.00
7	TYR	CZ	-40.63	-29.23	69.53	15.00
7	TYR	OH	-40.78	-30.58	69.62	15.00
7	TYR	C	-41.58	-22.91	69.65	15.00
7	TYR	0	-41.37	-22.64	70.84	15.00
8	ARG	N	-41.80	-21.97	68.74	15.00
8	ARG	CA	-41.78	-20.56	69.11	15.00
8	ARG	CB	-41.99	-19.67	67.87	15.00
8	ARG	CG	-40.88	-19.77	66.84	15.00
8	ARG	CD	-41.23	-19.01	65.57	15.00
8	ARG	NE	-42.50	-19.45	65.01	15.00
8	ARG	CZ	-42.96	-19.10	63.81	15.00
8	ARG	NH1	-42.24	-18.30	63.03	15.00
8	ARG	NH2	-44.15	-19.53	63.40	15.00
8	ARG		-42.85	-20.28	70.15	15.00
8	ARG	0	-42.57	-19.72	71.21	15.00
9	LYS	N	-44.06	-20.77	69.88	15.00
9	LYS		-45.18	-20.55	70.78	15.00
9	LYS		-46.50	-20.64	70.02	15.00
9	LYS		-46.63	-21.81	69.10	15.00
9	LYS	CD	-47.93	-21.71	68.33	15.00
	LYS		-48.22		67.54	15.00
9	LYS	NZ		-22.90	66.80	15.00
9	LYS	С	-45.22	-21.43	72.02	15.00
9	LYS	0	-46.25	-21.56	72.67	15.00
	LYS	N	-44.08	-22.02	72.38	15.00
	LYS		-43.97	-22.87	73.56	15.00
10	LYS		-43.66	-24.32	73.16	15.00
	LYS			-25.04	72.44	15.00
10	LYS	CD	-44.37	-26.46	72.08	15.00

10 LYS CE	-45.44	-27.16	71.27	15.00
10 LYS NZ	-45.80			15.00
10 LYS C	-42.90	-22.38	74.54	15.00
10 LYS O	-42.69	-22.99	75.59	15.00
11 GLY N	-42.19	-21.30	74.19	15.00
11 GLY CA	-41.15	-20.78	75.05	15.00
11 GLY C	-39.83	-21.52	74.90	15.00
11 GLY O	-38.95	-21.41	75.74	15.00
12 TYR N	-39.69	-22.25	73.79	15.00
12 TYR CA	-38.48	-23.03	73.51	15.00
12 TYR CB	-38.82	-24.26	72.67	15.00
12 TYR CG	-39.21		73.39	15.00
12 TYR CD1	-40.43	-25.65	74.06	15.00
12 TYR CE1	-40.85	-26.87	74.58	15.00
12 TYR CD2	-38.41	-26.68	73.27	15.00
12 TYR CE2	-38.83		73.79	15.00
12 TYR CZ	-40.05		74.44	15.00
12 TYR OH	-40.47	-29.19	74.93	15.00
12 TYR C	-37.45	-22.25	72.72	15.00
12 TYR O	-36.33	-22.72	72.54	15.00
13 VAL N	-37.85	-21.09	72.19	15.00
13 VAL CA	-36.94	-20.29	71.37	15.00
13 VAL CB	-37.40	-20.32	69.88	15.00
13 VAL CG1	-38.59	-19.43	69.67	15.00
13 VAL CG2	-36.26	-19.93	68.97	15.00
13 VAL C	-36.77	-18.86	71.87	15.00
13 VAL O	-37.69	-18.26	72.43	15.00
14 THR N	-35.55	-18.35	71.69	15.00
14 THR CA	-35.15	-17.00	72.11	15.00
14 THR CB	-33.72	-17.00	72.67	15.00
14 THR OG1	-32.83	-17.53	71.69	15.00
14 THR CG2	-33.64	-17.86	73.91	15.00
14 THR C	-35.21	-15.98	70.97	15.00
L4 THR O	-35.26	-16.36	69.81	15.00
L5 PRO N	-35.22	-14.68	71.31	15.00
L5 PRO CD	-35.25	-14.08	72.65	15.00
L5 PRO CA	-35.27	-13.64	70.28	15.00
15 PRO CB	-35.03	-12.35	71.08	15.00
.5 PRO CG	-34.46	-12.83	72.43	15.00
.5 PRO C	-34.23	-13.82	69.17	15.00
.5 PRO O	-33.14	-14.35	69.41	15.00
6 VAL N		-13.42	67.96	15.00
.6 VAL CA	-33.72	-13.53	€6.81	15.00

16 VAL CB	-34.45		65.50	15.00
16 VAL CG1	-33.63	-13.60	64.31	15.00
16 VAL CG2	-35.81	-13.78	65.47	15.00
16 VAL C	-32.52	-12.61	66.97	15.00
16 VAL O	-32.67	-11.42	67.25	15.00
17 LYS N	-31.32	-13.17	66.80	15.00
17 LYS CA	-30.09	-12.42	66.91	15.00
17 LYS CB	-29.08	-13.18	67.77	15.00
17 LYS CG	-29.01	-12.72	69.22	15.00
17 LYS CD	-30.27	-13.09	70.02	15.00
17 LYS CE	-30.18	-14.48	70.64	15.00
17 LYS NZ	-29.12	-14.54	71.68	15.00
17 LYS C	-29.49		65.54	15.00
17 LYS O	-29.82	-12.81	64.56	15.00
18 ASN N	-28.62	-11.15	65.49	15.00
18 ASN CA	-27.91	-10.77	64.27	15.00
18 ASN CB	-28.01	-9.26	64.05	15.00
18 ASN CG	-27.09		62.94	15.00
18 ASN OD1	-26.98	-9.42	61.89	15.00
18 ASN ND2	-26.38	-7.69	63.19	15.00
18 ASN C	-26.45	-11.16	64.43	15.00
18 ASN O	-25.79	-10.73	65.37	15.00
19 GLN N	-25.94	-11.96	63.51	15.00
19 GLN CA	-24.56	-12.40	63.60	15.00
19 GLN CB	-24.34	-13.65	62.77	15.00
19 GLN CG	-24.88	~13.58	61.37	15.00
19 GLN CD	-24.42	-14.73	60.53	15.00
19 GLN OE1	-25.17	-15.28	59.73	15.00
19 GLN NE2	-23.15	-15.09	60.68	15.00
19 GLN C	-23.52	-11.35	63.24	15.00
19 GLN 0	-22.35	-11.49	63.60	15.00
20 GLY N	-23.94	-10.31	62.53	15.00
20 GLY CA	-23.02	-9.26	62.14	15.00
20 GLY C	-22.10	-9.67	61.00	15.00
20 GLY O	-22.51	-10.40	60.11	15.00
21 GLN N	-20.85	-9.20	61.04	15.00
21 GLN CA	-19.86	-9.50	60.01	15.00
21 GLN CB	-18.92	-8.30	59.78	15.00
21 GLN CG	-19.60	-6.94	59.55	15.00
21 GLN CD	-20.68	-6.99	58.49	15.00
21 GLN OE1	-20.49	-7.53	57.41	15.00
21 GLN NE2	-21.85	-6.43	58.81	15.00
21 GLN C	-19.03	-10.74	60.38	15.00

21	GLN	0	-18.02	-11.06	59.74	15.00
22	CYS	N	-19.41	-11.42	61.44	15.00
22	CYS	CA	-18.69	-12.60	61.88	15.00
22	CYS	C	-19.37	-13.86	61.33	15.00
22	CYS	0	-20.59	-13.94	61.27	15.00
22	CYS	CB	-18.63	-12.61	63.41	15.00
22	CYS	SG	-17.84	-14.05	64.16	15.00
23	GLY	N	-18.58	-14.81	60.84	15.00
23	GLY	CA	-19.15	-16.03	60.30	15.00
23	GLY	C	-19.41	-17.03	61.41	15.00
23	GLY	0	-18.90	-18.15	61.38	15.00
24	SER	N	-20.22	-16.62	62.37	15.00
24	SER	CA	-20.56	-17.43	63.53	<b>i5.00</b>
24	SER	CB	-20.36	-16.59	64.79	15.00
24	SER	OG	-21.14	-15.41	64.68	15.00
24	SER	C	-21.99	-17.95	63.49	15.00
24	SER	0	-22.60	-18.20	64.53	15.00
25	CYS	N	-22.55	-18.14	62.30	15.00
25	CYS	CA	-23.91	-18.65	62.20	15.00
25	CYS	CB	-24.41	-18.63	60.75	15.00
25	CYS	SG	-23.48	-19.60	59.54	15.00
25	CYS	C	-23.96	-20.05	62.82	15.00
25	CYS	0	-24.97	-20.45	63.40	15.00
25	INH	C1	-27.24	-9.28	57.72	15.00
25	INH	C2	-26.55	-9.60	58.90	15.00
25	INH	C3	-25.31	-10.22	58.84	15.00
25	INH	C4	-24.73	-10.54	57.61	15.00
25	INH	C5	-25.43	-10.21	56.44	15.00
25	INH	C6	-26.67	-9.59	56.49	15.00
25	INH	C7	-23.41	-11.26	57.54	15.00
25	INH	08	-23.43	-12.63	57.98	15.00
25	INH	C9	-22.90	-13.56	57.08	15.00
25	INH	010	-21.75	-13.43	56.65	15.00
25	INH	C11	-23.40	-15.62	55.77	15.00
25	INH	C12	-22.32	-15.20	54.77	15.00
25	INH	C13	-22.79	-14.65	53.42	15.00
25	INH	C14	-21.66	-14.80	52.41	15.00
25	INH	C15	-24.07	-15.33	52.91	15.00
25	INH	C16	-23.13	-17.06	56.23	15.00
25	INH	017	-23.79	-17.98	55.74	15.00
25	INH	N18			57.12	
25	INH	C19			57.60	
25	INH	N20			56.73	

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25		-21.96	-18.81	59.10	15.00
25		-21.89	-17.72	59.66	15.00
25		-19.42	-29.07	54.27	15.00
25		-20.11	-28.05	54.90	15.00
25		-19.45	-26.88	55.23	15.00
25		-18.09	-26.70	54.93	15.00
25	INH C27	-17.41	-27.74	54.30	15.00
25	INH C28	-18.06	-28.92	53.96	15.00
25	INH C29	-17.39	-25.41	55.26	15.00
25	INH 030	-18.05	-24.15	55.06	15.00
25	INH C31	-19.20	-23.80	55.81	15.00
25	INH 032	-20.33	-23.84	55.32	15.00
25	INH C33	-20.15	-23.05	57.92	15.00
25	INH C34	-20.47	-24.13	58.95	15.00
25	INH C35	-21.49	-25.18	58.56	15.00
25	INH C36	-22.36	-24.69	57.40	15.00
25	INH C37	-20.73	-26.42	58.16	15.00
25	INH C38	-19.89	-21.74	58.63	15.00
25	INH 039	-18.75	-21.39	58.90	15.00
25	INH N40	-20.97	-21.01	58.95	15.00
25	INH C41	-20.91	-19.72	59.64	15.00
25	INH N42	-19.01	-23.44	57.08	15.00
26	TRP N	-22.84	-20.77	62.76	15.00
26	TRP CA	-22.76	-22.11	63.33	15.00
26	TRP CB	-21.47	-22.81	62.88	15.00
26	TRP CG	-20.24	-22.16	63.40	15.00
26	TRP CD2	-19.56	-22.46	64.62	15.00
26	TRP CE2	-18.51	-21.53	64.75	15.00
26	TRP CE3	-19.74	-23.42	65.63	15.00
26	TRP CD1	-19.59	-21.11	62.84	15.00
26	TRP NE1	-18.55	-20.72	63.65	15.00
26	TRP CZ2	-17.64	-21.52	65.85	15.00
26	TRP CZ3	-18.88	-23.42	66.72	15.00
26	TRP CH2	-17.84	-22.47	66.82	15.00
26	TRP C	-22.82	-22.02	64.87	15.00
26	TRP O	-23.31	-22.93	65.53	15.00
27	ALA N	-22.34	-20.92	65.44	15.00
27	ALA CA	-22.36	-20.73	66.89	15.00
27	ALA CB	-21.43	-19.61	67.30	15.00
27	ALA C	-23.79	-20.43	673.2	15.00
27	ALA O	-24.29	-21.00	68.29	15.00
28	PHE N	24.48	-19.57	66.58	15.00
28	PHE CA	-25.85	-19.25	66.92	15.00

28 PHE CB	-26.38	-18.14	66.01	15.00
28 PHE CG	-25.87	-16.78	66.39	15.00
28 PHE CD1	-24.63	-16.33	65.94	15.00
28 PHE CD2	-26.61	-15.96	67.22	15.00
28 PHE CE1	-24.14	-15.10	66.33	15.00
28 PHE CE2	-26.11	-14.72	67.62	15.00
28 PHE CZ	-24.88	-14.29	67.17	15.00
28 PHE C	-26.73	-20.49	66.83	15.00
28 PHE O	-27.48	-20.80	67.75	15.00
29 SER N	-26.60	-21.24	65.74	15.00
29 SER CA	-27.36	-22.46	65.54	15.00
29 SER CB	-26.91	-23.14	64.25	15.00
29 SER OG	-27.55	-24.39	64.08	15.00
29 SER C	-27.17	-23.43	66.70	15.00
29 SER O	-28.14	-23.89	67.30	15.00
30 SER N	-25.91	-23.75	67.01	15.00
30 SER CA	-25.57	-24.67	68.09	15.00
30 SER CB	-24.06	-24.68	68.33	15.00
30 SER OG	-23.33	-25.06	67.19	15.00
30 SER C	-26.28	-24.24	69.38	15.00
30 SER O	-27.01	-25.02	70.01	15.00
31 VAL N	-26.09	-22.97	69.73	15.00
31 VAL CA	-26.67	-22.39	70.93	15.00
31 VAL CB	-26.13	-20.95	71.14	15.00
31 VAL CG1	-27.14	-20.07	71.79	15.00
31 VAL CG2	-24.87	-20.99	71.99	15.00
31 VAL C	-28.21	-22.46	70.89	15.00
31 VAL O	-28.86	-22.68	71.92	15.00
32 GLY N	-28.79	-22.36	69.70	15.00
32 GLY CA	-30.23	-22.42	69.58	15.00
32 GLY C	-30.77	-23.77	69.99	15.00
32 GLY 0	-31.84	-23.88	70.58	15.00
33 ALA N	-30.04	-24.83	69.66	15.00
33 ALA CA		-26.18	70.01	15.00
33 ALA CB	-29.67	-27.20	69.23	15.00
33 ALA C	-30.27	-26.36	71.50	15.00
33 ALA O	-31.10	-26.99	72.17	15.00
34 LEU N	-29.20	-25.80	72.04	15.00
34 LEU CA	-28.91	-25.88	73.47	15.00
34 LEU CB	-27.55	-25.25	73.77	15.00
4 LEU CG		-26.14	73.47	15.00
4 LEU CD1		-25.37	73.60	15.00
4 LEU CD2	-26.34	-27.30	74.42	15.00

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34 LEU C	-30.00	-25.20	74.29	15.00
34 LEU O	-30.37	-25.66	75.38	15.00
35 GLU N	-30.56	-24.10	73.78	15.00
35 GLU CA	-31.61	-23.39	74.49	15.00
35 GLU CB	-31.82	-22.00	73.88	15.00
35 GLU CG	-30.62	-21.08	74.05	15.00
35 GLU CD	-30.60	-19.92	73.08	15.00
35 GLU OE1	-31.49	-19.83	72.21	15.00
35 GLU OE2	-29.66	-19.10	73.17	15.00
35 GLU C	-32.91	-24.18	74.47	15.00
35 GLU O	-33.62	-24.26	75.47	15.00
36 GLY N	-33.21	-24.80	73.33	15.00
36 GLY CA	-34.43	-25.58	73.22	15.00
36 GLY C	-34.49	-26.72	74.22	15.00
36 GLY O	<del>-</del> 35.52	-26.94	74.86	15.00
37 GLN N	-33.38	-27.43	74.36	15.00
37 GLN CA	-33.27	-28.55	75.29	15.00
37 GLN CB	-31.99	-29.34	75.02	15.00
37 GLN CG	-32.04	-30.08	73.69	15.00
37 GLN CD	-33.27	-30.95	73.58	15.00
37 GLN OE1	-33.55	-31.77	74.45	15.00
37 GLN NE2	-34.04	-30.77	72.52	15.00
37 GLN C	-33.31	-28.07	76.74	15.00
37 GLN O	-33.94	-28.69	77.60	15.00
38 LEU N	-32.66	-26.94	77.00	15.00
38 LEU CA	-32.66	-26.34	78.34	15.00
38 LEU CB	-31.99	-24.97	78.30	15.00
38 LEU CG	-31.70	-24.28	79.63	15.00
38 LEU CD1	-30.58	-25.02	80.35	15.00
38 LEU CD2	-31.32	-22.84	79.40	15.00
38 LEU C	-34.11	-26.19	78.79	15.00
38 LEU O	-34.52	-26.71	79.82	15.00
39 LYS N	-34.91	-25.51	77.97	15.00
39 LYS CA	-36.31		78.27	15.00
39 LYS CB	-36.98	-24.53	77.12	15.00
39 LYS CG	-38.48	-24.37	77.25	15.00
39 LYS CD	-38.89	-23.62	78.49	15.00
39 LYS CE	-40.38	-23.44	78.51	15.00
39 LYS NZ		-22.84	79.78	15.00
39 LYS C		-26.52	78.50	15.00
39 LYS O	-37.78	-26.76	79.44	15.00
40 LYS N		-27.62	77.68	15.00
40 LYS CA	-37.32	-2893	77.83	15.00

40	LYS	CB	-36.78	-29.89	76.76	15.00
40	LYS	CG	-37.56	-31.18	76.57	15.00
40	LYS	CD	-36.89	-32.09	75.54	15.00
40	LYS	CE	-37.73	-33.32	75.24	15.00
40	LYS	NZ	-39.05	-32.97	74.63	15.00
40	LYS	С	-37.07	-29.52	79.22	15.00
40	LYS	0	-38.00	-29.92	79.93	15.00
41	LYS	N	-35.80	-29.51	79.64	15.00
41	LYS	CA	-35.41	-30.07	80.92	15.00
41	LYS	CB	-33.92	-30.40	80.91	15.00
41	LYS	CG	-33.48	-31.19	79.67	15.00
41	LYS	CD	-34.36	-32.41	79.42	15.00
41	LYS	CE	-34.05	-33.08	78.08	15.00
41	LYS	NZ	-34.99	-34.21	77.78	15.00
41	LYS	С	-35.76	-29.21	82.13	15.00
41	LYS	0	-36.58	-29.61	82.96	15.00
42	THR	N	-35.17	-28.02	82.25	15.00
42	THR	CA	-35.41	-27.14	83.39	15.00
42	THR	CB	-34.27	-26.12	83.52	15.00
42	THR	OG1	-34.29	-25.25	82.38	15.00
42	THR	CG2	-32.94	-26.82	83.57	15.00
42	THR	С	-36.72	-26.37	83.39	15.00
42	THR	0	-37.10	-25.79	84.41	15.00
43	GLY	N	-37.40	-26.33	82.25	15.00
43	GLY	CA	-38.65	-25.60	82.15	15.00
43	GLY	С	-38.47	-24.08	82.03	15.00
43	GLY	0	-39.43	-23.33	81.98	15.00
44	LYS		-37.24	-23.57	82.01	15.00
44	LYS		-37.05	-22.13	81.89	15.00
44	LYS	CB	-36.55	-21.53	83.20	15.00
	LYS	CG	-37.49	-21.67	84.38	15.00
44	LYS		-36.91	-20.99	85.61	15.00
	LYS		-35.68	-21.70	86.11	15.00
44	LYS	NZ	-36.03	-23.03	86.65	15.00
	LYS		-36.05	-21.83	80.80	15.00
44	LYS	0	-35.00	-22.48	80.70	15.00
45	LEU		-36.39	-20.84	79.97	15.00
	LEU		-35.55	-20.41	78.86	15.00
	LEU		-36.43	-19.88	77.73	15.00
	LEU		-35.82	-19.71	76.33	15.00
45			-35.62	-21.08	75.69	15.00
	LEU		-36.74	-1.8.87	75.48	15.00
45	LEU	С	-34.58	-19.34	79.32	15.00

			***	
45 LEU O	-34.92		80.13	15.00
46 LEU N	-33.36		78.80	15.00
46 LEU CA	-32.32	-18.44	79.15	15.00
46 LEU CB	-31.47		80.28	15.00
46 LEU CG	-30.58	-18.14	81.17	15.00
46 LEU CD1	-29.23	-17.96	80.55	15.00
46 LEU CD2	-31.23	-16.81	81.44	15.00
46 LEU C	-31.51	-18.26	77.86	15.00
46 LEU O	-31.48	-19.15	77.01	15.00
47 ASN N	-30.92	-17.08	77.67	15.00
47 ASN CA	-30.14	-16.82	76.47	15.00
47 ASN CB	-30.22	-15.35	76.06	15.00
47 ASN CG	-31.33	-15.09	75.06	15.00
47 ASN OD1	-32.44	-14.74	75.43	15.00
47 ASN ND2	-31.04	-15.28	73.78	15.00
47 ASN C	-28.69	-17.26	76.64	15.00
47 ASN 0	-27.98	-16.75	77.50	15.00
48 LEU N	-28.27	-18.21	75.82	15.00
48 LEU CA	-26.92	-18.73	75.89	15.00
48 LEU CB	-26.89	-20.21	75.51	15.00
48 LEU CG	-27.53	-21.15	76.55	15.00
48 LEU CD1	-27.34	-22.60	76.17	15.00
48 LEU CD2	-26.88	-20.92	77.88	15.00
48 LEU C	-25.93	-17.89	75.07	15.00
48 LEU O	-26.32	-17.11	74.20	15.00
49 SER N	-24.64	-18.08	75.35	15.00
49 SER CA	-23.56	-17.34	74.70	15.00
49 SER CB	-22.47	-17.07	75.75	15.00
49 SER OG	-21.31	-16.50	75.18	15.00
49 SER C	-22.92	-17.91	73.43	15.00
49 SER O	-22.16	-18.88	73.48	15.00
50 PRO N	-23.22	-17.30	72.26	15.00
50 PRO CD	-24.28	-16.31	72.02	15.00
50 PRO CA	-22.65	-17.75	70.98	15.00
50 PRO CB	-23.42	-16.92	69.95	15.00
50 PRO CG	-24.70	-16.64	70.62	15.00
50 PRO C	-21.16	-17.38	70.95	15.00
0 PRO O	-20.34	-18.08	70.35	15.00
I GLN N	-20.81	-16.27	71.59	15.00
1 GLN CA		-15.80	71.65	15.00
1 GLN CB		-14.43	72.34	15.00
1 GLN CG	-17.95	-13.79	72.39	15.00
1 GLN CD		-13.23	71.05	15.00
			( ± . U 🗸	<b>10.00</b>

51	GLN OE1	-18.21	-12.62	70.30	15.00
51	GLN NE2	-16.19	-13.42	70.77	15.00
51	GLN C	-18.58	-16.83	72.38	15.00
51	GLN O	-17.46	-17.15	71.96	15.00
52	ASN N	-19.13	-17.42	73.44	15.00
52	ASN CA	-18.41	-18.42	74.24	15.00
52	ASN CB	-19.31	-18.96	75.35	15.00
52	ASN CG	-18.59	-19.90	76.31	15.00
52	ASN OD1	-19.23	-20.62	77.07	15.00
52	ASN ND2	-17.27	-19.89	76.28	15.00
52	ASN C	-17.91	-19.55	73.35	15.00
52	asn o	-16.84	-20.12	73.59	15.00
53	LEU N	-18.67	-19.86	72.31	15.00
53	LEU CA	-18.33	-20.90	71.35	15.00
53	LEU CB	-19.58	-21.37	70.60	15.00
53	LEU CG	-20.63	-22.10	71.42	15.00
53	LEU CD1	-21.81	-22.44	70.55	15.00
53	LEU CD2	-20.01	-23.36	72.01	15.00
53	LEU C	-17.31	-20.42	70.34	15.00
53	LEU O	-16.37	-21.13	69.99	15.00
54	VAL N	-17.52	-19.19	69.87	15.00
54	VAL CA	-16.66	-18.57	68.87	15.00
	VAL CB	-17.16	-17.14	68.56	15.00
	VAL CG1	-16.22	-16.45	67.59	15.00
	VAL CG2	-18.57	-17.20	68.00	15.00
	VAL C	-15.20	-18.52	69.30	15.00
54	VAL O	-14.31	-18.88	68.53	15.00
55	ASP N	-14.96	-18.09	70.54	15.00
55	ASP CA	-13.61	-17.98	71.07	15.00
55	ASP CB	-13.58	-16.93	72.18	15.00
55	ASP CG	-14.14	-15.59	71.76	15.00
	ASP OD1	-14.26	-15.33	70.55	15.00
	ASP OD2	-14.45	-14.78	72.67	15.00
	ASP C		-19.26	71.65	15.00
	ASP O	-11.80	-19.39	71.72	15.00
	CYS N	-13.86	-20.19	72.09	15.00
	CYS CA	-13.36	-21.41	72.73	15.00
	CYS C	-13.29	-22.71	71.94	15.00
	CYS O	-12.43	-23.55	72.20	15.00
	CYS CB	-14.10	-21.61	74.04	15.00
	CYS SG	-14.21	-20.10	75.06	15.00
	VAL N	-14.17	-22.93	70.96	15.00
57	VAL CA	-14.13	-24.1.7	70.18	15.00

57 VAL CB	-15.44	-24.43	69.39	15.00
57 VAL CG1	-15.38	-25.80	68.73	15.00
57 VAL CG2	-16.63	-24.33	70.30	15.00
57 VAL C	-12.97	-24.10	69.19	15.00
57 VAL O	-13.17	-23.81	68.02	15.00
58 SER N	-11.76	-24.40	69.66	15.00
58 SER CA	-10.57	-24.35	68.82	15.00
58 SER CB	-9.34	-24.73	69.63	15.00
58 SER OG	-9.44	-26.08	70.07	15.00
58 SER C	-10.65	-25.25	67.61	15.00
58 SER O	-9.90	-25.09	66.65	15.00
59 GLU N	-11.53	-26.25	67.65	15.00
59 GLU CA	-11.69	-27.16	66.53	15.00
59 GLU CB	-12.57	-28.35	66.92	15.00
59 GLU CG	-12.00	-29.23	68.00	15.00
59 GLU CD	-11.98	-28.56	69.36	15.00
59 GLU OE1	-13.00	-27.95	69.73	15.00
59 GLU OE2	-10.95	-28.66	70.06	15.00
59 GLU C	-12.30	-26.41	65.35	15.00
59 GLU O	-12.28	-26.89	64.22	15.00
60 ASN N	-12.89	-25.26	65.63	15.00
60 ASN CA	-13.53	-24.45	64.61	15.00
60 ASN CB	-14.95	-24.06	65.03	15.00
60 ASN CG	-15.92	-25.21	64.90	15.00
60 ASN OD1	-17.09	-25.08	65.23	15.00
60 ASN ND2	-15.45	-26.34	64.38	15.00
60 ASN C	-12.71	-23.22	64.26	15.00
60 ASN O	-11.73	-22.92	64.93	15.00
61 ASP N	-13.13	-22.50	63.23	15.00
61 ASP CA	-12.41	-21.34	62.75	15.00
61 ASP CB	-12.53	-21.28	61.22	15.00
61 ASP CG	-11.20	-21.01	60.53	15.00
61 ASP OD1	-10.34	-20.32	61.12	15.00
61 ASP OD2	-11.03	-21.49	59.40	15.00
61 ASP C	-12.80	-19.99	63.35	15.00
61 ASP 0	-12.23	-18.95	62.99	15.00
62 GLY N	-13.75	-19.97	64.28	15.00
62 GLY CA	-14.17	-18.71	64.87	15.00
62 GLY C	-15.12	-18.03	63.91	15.00
62 GLY O	-16.13	-18.61	63.51	15.00
63 CYS N	-14.80	-16.81	63.48	15.00
63 CYS CA	-15.65	-16.09	62.52	15.00
63 CYS C	-15.45	-16.63	61.12	

63 CYS (	-16.10	-16.16	60.18	15.00
63 CYS (	CB ~15.34	-14.59	62.52	15.00
63 CYS	SG -15.84	-13.72	64.03	15.00
64 GLY 1	N -14.52	-17.56	60.95	15.00
64 GLY	CA -14.26	-18.14	59.65	15.00
64 GLY	-15.17	-19.33	59.40	15.00
64 GLY	-15.50	-19.67	58.26	15.00
65 GLY 1	· -15.60	-19.99	60.48	15.00
65 GLY C	CA -16.47	-21.14	60.33	15.00
65 GLY C	-16.39	-22.20	61.41	15.00
65 GLY C	-15.56	-22.15	62.31	15.00
66 GLY N	-17.28	-23.18	61.30	15.00
66 GLY C	A -17.31	-24.25	62.27	15.00
66 GLY C	-18.50	-25.16	62.14	15.00
66 GLY C	-19.48	-24.83	61.48	15.00
67 TYR N	-18.43	-26.32	62.78	15.00
67 TYR C	A -19.51	-27.30	62.74	15.00
67 TYR C	B -18.97	-28.70	62.45	15.00
67 TYR C	G -18.28	-28.86	61.12	15.00
67 TYR C	D1 -19.02	-29.00	59.94	15.00
67 TYR C	E1 -18.40	-29.23	58.72	15.00
67 TYR C			61.04	15.00
67 TYR C	E2 -16.26	-29.16	59.82	15.00
67 TYR C	<b>z</b> -17.02	-29.31	58.67	15.00
67 TYR C		-29.51	57.46	15.00
67 TYR C		-27.29	64.10	15.00
67 TYR O		-27.14	65.12	15.00
68 MET N	-21.48	-27.47	64.12	15.00
68 MET C		-27.48	65.36	15.00
68 MET C		-27.60	65.07	15.00
68 MET C			64.29	15.00
68 MET S			62.55	15.00
68 MET C			61.83	15.00
68 MET C		-28.62	66.27	15.00
68 MET O			67.47	15.00
69 THR N		-29.79	65.68	15.00
69 THR C			66.43	15.00
69 THR C		-32.18	65.50	15.00
69 THR O		-31.81	64.42	15.00
69 THR C	G2 -22.21	-32.71	64.94	15.00
69 THR C	-19.90		67.25	15.00
69 THR O			68.41	15.00
70 ASN N	-18.98	-29.85	66.69	15.00

70 ASN CA	-17.74		67.37	15.00
70 ASN CB	-16.79	-	66.42	15.00
70 ASN CG	-16.05	-29.71	65.50	15.00
70 ASN OD1	-14.84	<del>-</del>	65.39	15.00
70 ASN ND2	-16.78		64.83	15.00
70 ASN C	-18.01	-28.62	68.59	15.00
70 ASN 0	-17.32	-28.74	69.60	15.00
71 ALA N	-19.03		68.48	15.00
71 ALA CA	-19.43		69.56	15.00
71 ALA CB	-20.37	-25.81	69.04	15.00
71 ALA C	-20.12	-27.68	70.67	15.00
71 ALA O	-20.03	-27.32	71.84	15.00
72 PHE N	-20.85	-28.72	70.28	15.00
72 PHE CA	-21.54	-29.59	71.22	15.00
72 PHE CB	-22.36	-30.65	70.47	15.00
72 PHE CG	-23.54	-30.11	69.74	15.00
72 PHE CD1	-24.28	-29.06	70.24	15.00
72 PHE CD2	-23.91	-30.66	68.53	15.00
72 PHE CE1	-25.36	-28.56	69.55	15.00
72 PHE CE2	-25.00	-30.17	67.83	15.00
72 PHE CZ	-25.72	-29.12	68.34	15.00
72 PHE C	-20.46	-30.29	72.05	15.00
72 PHE O	-20.43	-30.18	73.28	15.00
73 GLN N	-19.58	-30.98	71.34	15.00
73 GLN CA	-18.47	-31.71	71.92	15.00
73 GLN CB	-17.53	-32.13	70.78	15.00
73 GLN CG	-16.60	-33.30	71.06	15.00
73 GLN CD	-17.31	-34.63	71.03	15.00
73 GLN OE1	-17.80	-35.13	72.06	15.00
73 GLN NE2	-17.35	-35.25	69.85	15.00
73 GLN C	-17.74	-30.81	72.93	15.00
73 GLN 0	-17.27	-31.26	73.97	15.00
74 TYR N	-17.66	-29.51	72.63	15.00
74 TYR CA	-17.00	-28.57	73.51	15.00
74 TYR CB	-16.75	-27.21	72.81	15.00
74 TYR CG	-16.41	-26.08	73.77	15.00
74 TYR CD1	-15.19	-26.05	74.43	15.00
74 TYR CE1	-14.91	-25.06	75.37	15.00
74 TYR CD2	-17.34	-25.09	74.06	15.00
74 TYR CE2	-17.07	-24.10	75.00	15.00
74 TYR CZ	-15.86	-24.09	75.66	15.00
74 TYR OH	-15.60	-23.15	76.63	15.00
74 TYR C	-17.75	-28.36	74.82	15.00

74	TYR	. 0	-17.14	-28.37	75.89	15.00
75	VAL	N	-19.07	-28.16	74.78	15.00
75	VAL	CA	-19.85	-27.94	76.02	15.00
75	VAL	CB	-21.30	-27.45	75.73	15.00
75	VAL	CG1	-22.06	-27.23	77.04	15.00
75	VAL	CG2	-21.27	-26.15	74.92	15.00
75	VAL	С	-19.87	-29.20	76.91	15.00
75	VAL	0	-20.07	-29.13	78.12	15.00
76	GLN	N	-19.65	-30.35	76.29	15.00
76	GLN	CA	-19.62	-31.60	77.01	15.00
76	GLN	CB	-19.93	-32.73	76.04	15.00
76	GLN	CG	-20.02	-34.11	76.67	15.00
76	GLN	CD	-19.74	-35.20	75.68	15.00
76	GLN	OE1	-20.14	-35.11	74.52	15.00
76	GLN	NE2	-19.01	-36.22	76.11	15.00
76	GLN	C	-18.25	-31.81	77.66	15.00
76	GLN	0	-18.14	-32.17	78.85	15.00
77	LYS	N	-17.18	-31.59	76.89	15.00
77	LYS	CA	-15.82	-31.78	77.39	15.00
77	LYS	CB	-14.83	-31.98	76.24	15.00
77	LYS	CG	-14.56	-30.75	75.41	15.00
77	LYS	CD	-13.69	-31.09	74.19	15.00
77	LYS	CE	-14.41	-32.05	73.24	15.00
77	LYS		-13.62	-32.45	72.02	15.00
77	LYS		-15.34	-30.65	78.29	15.00
77	LYS	0	-14.37	-30.81	79.01	15.00
	ASN	N	-15.97	-29.49	78.22	15.00
78	ASN		-15.60	-28.37	79.07	15.00
	ASN		-15.58	-27.07	78.27	15.00
	ASN		-15.38	-25.85	79.16	15.00
	ASN		-14.28	-25.58	79.64	15.00
	ASN		-16.45	-25.11	79.37	15.00
	ASN		-16.62	-28.28	80.20	15.00
	ASN			-27.67	81.23	15.00
	ARG		-17.77	-28.91	79.97	15.00
79	ARG		-18.89	-28.98	80.92	15.00
79	ARG		-18.58	-29.92	82.11	15.00
79	ARG		-17.47	-29.49	83.06	15.00
79	ARG		-16.62	-30.66	83.51	15.00
	ARG		-17.44	-31.76	84.00	15.00
79				-32.75	83.23	15.00
	ARG			-32.77	81.93	15.00
79	ARG	NH2	-18.73	-33.66	83.74	15.00

79	AR(	G C	-19.45	-27.65	81.39	15.00
79	AR	GO	-19.42	-27.31	82.57	15.00
80	GL:	Y N	-20.00	-26.92	80.44	15.00
80	GL;	CA	-20.58	-25.63	80.75	15.00
80	GL1	C	-20.41	-24.62	79.63	15.00
80	GL3	0	-19.39	-24.61	78.93	15.00
81	ILE	e n	-21.44	-23.82	79.44	15.00
81	ILE	E CA	-21.45	-22.77	78.45	15.00
81	ILE	CB	-22.27	-23.14	77.21	15.00
81	ILE	CG2	-23.71	-23.46	77.59	15.00
81	ILE	CG1	-22.18	-22.01	76.17	15.00
81	ILE	CD1	-22.74	-22.34	74.79	15.00
81	ILE	c c	-22.09	-21.59	79.18	15.00
81	ILE	0	-23.08	-21.77	79.89	15.00
82	ASP	N	-21.48	-20.42	79.06	15.00
82	ASP	CA	-21.99	-19.22	79.72	15.00
82	ASP	CB	-20.95	-18.10	79.68	15.00
82	ASP	CG	-19.75	-18.37	80.56	15.00
82	ASP	OD1	-18.66	-17.89	80.20	15.00
82	ASP	OD2	-19.89	-19.03	81.62	15.00
82	ASP	C	-23.27	-18.70	79.09	15.00
82	ASP	0	-23.67	-19.11	78.01	15.00
83	SER	N	-23.91	-17.78	79.79	15.00
83	SER	CA	-25.12	-17.15	79.30	15.00
83	SER	CB	-26.03	-16.79	80.46	15.00
83	SER	OG	-25.32	-16.05	81.44	15.00
83	SER	С	-24.66	-15.90	78.58	15.00
83	SER	0	-23.49	-15.51	78.67	15.00
84	GLU	N	-25.57	-15.24	77.87	15.00
84	GLU	CA	-25.22	-14.02	77.16	15.00
84	GLU	CB	-26.40	-13.50	76.35	15.00
84	GLU		-26.09	-12.30	75.46	15.00
	GLU		-25.06	-12.58	74.36	15.00
		OE1	-25.03	-13.69	73.78	15.00
		OE2	-24.28		74.06	15.00
	GLU		-24.70	-12.95	78.14	15.00
84	GLU	0	-23.64	-12.37	77.92	15.00
	ASP		-25.41	-12.69	79.23	15.00
	ASP		-24.95	-11.70	80.20	15.00
	ASP		-25.99	-11.46	81.31	15.00
25	ASP	CG	-25.59	-10.32	82.30	15.00
	ASP		-26.24	-10.19		15.00
85	ASP	OD2	-24.66	-9.54		

85	ASP	С	-23.63	-12.15	80.82	15.00
85	ASP	0	-22.86	-11.34	81.34	15.00
86	ALA	N	-23.31	-13.44	80.74	15.00
86	ALA	CA	-22.07	-13.91	81.32	15.00
86	ALA	CB	-22.24	-15.30	81.86	15.00
86	ALA	С	-20.90	-13.85	80.35	15.00
86	ALA	0	-19.74	-13.84	80.76	15.00
87	TYR	N	-21.19	-13.82	79.05	15.00
87	TYR	CA	-20.14	-13.77	78.03	15.00
87	TYR	CB	-19.67	-15.20	77.77	15.00
87	TYR	CG	-18.31	-15.36	77.13	15.00
87	TYR	CD1	-17.86	-14.49	76.13	15.00
87	TYR	CE1	-16.64	-14.70	75.51	15.00
87	TYR	CD2	-17.49	-16.43	77.48	15.00
87	TYR	CE2	-16.27	-16.64	76.86	15.00
87	TYR	CZ	-15.85	-15.78	75.88	15.00
87	TYR	OH	-14.65	-16.03	75.26	15.00
87	TYR	C	-20.73	-13.17	76.75	15.00
87	TYR	0	-20.92	-13.88	75.76	15.00
88	PRO	N	-20.99	-11.84	76.74	15.00
88	PRO	CD	-20.79	-10.92	77.87	15.00
88	PRO	CA	-21.57	-11.12	75.60	15.00
88	PRO	CB	-21.52	-9.66	76.07	15.00
88	PRO	CG	-21.75	-9.80	77.54	15.00
88	PRO	С	-20.91	-11.29	74.24	15.00
88	PRO	0	-19.71	-11.56	74.13	15.00
89	TYR	N	-21.72	-11.10	73.21	15.00
89	TYR	CA	-21.30	-11.23	71.83	15.00
89	TYR	CB	-22.51	-11.59	70.96	15.00
89	TYR	CG	-22.16	-12.03	69.56	15.00
89	TYR	CD1	-21.27	-13.09	69.35	15.00
89	TYR	CE1	-20.90	-13.46	68.07	15.00
89	TYR	CD2	-22.67	-11.37	68.45	15.00
89	TYR	CE2	-22.31	-11.74	67.17	15.00
89	TYR	CZ	-21.42	-12.78	66.98	15.00
89	TYR	OH	-21.02	-13.12	65.72	15.00
89	TYR	С	-20.67	-9.90	71.38	15.00
89	TYR	0	-21.33	-8.86	71.36	15.00
90	VAL	N	-19.38	-9.93	71.05	15.00
90	VAL	CA	-18.66	-8.73	70.61	15.00
90	VAL	CB	-17.25	-8.65	71.22	15.00
90	VAL	CG1	-17.32	-8.67	72.73	15.00
90	VAJ	CG2	-16.37	-9.78	70.68	15.00

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	VAL C	-18.54	-8.62	69.11	15.00
90	VAL O	-17.85	-7.75	68.61	15.00
91	GLY N	-19.14	-9.55	68.39	15.00
91	GLY CA	-19.10	-9.49	66.94	15.00
91	GLY C	-17.76	-9.77	66.31	15.00
91		-17.52	-9.39	65.15	15.00
92	GLN N	-16.88	-10.44	67.03	15.00
92	GLN CA	-15.56	-10.77	66.50	15.00
92	GLN CB	-14.74	-9.51	66.31	15.00
92	GLN CG	-14.49	-8.76	67.60	15.00
92	GLN CD	-13.93	-7.40	67.35	15.00
92	GLN OE	1 -12.83	-7.06	67.80	15.00
92	GLN NE	2 -14.67	-6.59	66.60	15.00
92	GLN C	-14.81	-11.73	67.41	15.00
92	GLN O	-15.19	-11.95	68.57	15.00
93	GLU N	-13.72	-12.26	66.89	15.00
93		-12.90	-13.22	67.59	15.00
93	GLU CB	-12.01	-13.95	66.59	15.00
93	GLU CG	-12.77	-14.60	65.42	15.00
93	GLU CD	-11.84	-15.07	64.31	15.00
93	GLU OE	L -10.67	-15.37	64.62	15.00
93	GLU OE	2 -12.27	-15.13	63.13	15.00
93	GLU C	-12.05	-12.56	68.66	15.00
93	GLU O	-11.53	-11.46	68.49	15.00
94	GLU N	-11.92	-13.25	69.78	15.00
94	GLU CA	-11.13	-12.78	70.91	15.00
94	GLU CB	-11.93	-11.79	71.76	15.00
94	GLU CG	-13.39	-12.15	71.95	15.00
94	GLU CD	-14.00	-11.50	73.18	15.00
94	GLU OE		-10.26	73.18	15.00
94	GLU OE2		-12.24	74.14	15.00
94	GLU C	-10.73	-13.98	71.74	15.00
	GLU O	-11.15	-15.10	71.47	15.00
	SER N	-9.88	-13.75	72.73	15.00
	SER CA	-9.40	-14.80	73.61	15.00
95	SER CB	-8.39	-14.21	74.60	15.00
95	SER OG	-7.22	-13.77	73.94	15.00
95	SER C	-10.57	-15.44	74.35	15.00
95		-11.48	-14.73	74.82	15.00
96		-10.55	-16.77	74.42	15.00
	CYS CA	-11.59	-17.52	75.13	15.00
	CYS C	-11.61	-17.01	76.56	15.00
96	CYS O	-10.57	-16.90	77.21	15.00

96 CYS CB	-11.31	-19.03	75.08	15.00
96 CYS SG	-12.44		76.07	15.00
97 MET N	-12.80		77.03	15.00
97 MET CA	-12.96		78.36	15.00
97 MET CB	-13.24		78.26	15.00
97 MET CG	-12.05		77.73	15.00
97 MET SD	-12.56	-12.29	76.76	15.00
97 MET CE	-13.48	-11.34	78.03	15.00
97 MET C	-14.07		79.14	15.00
97 MET O	-14.87		79.82	15.00
98 TYR N	-14.13	-18.10	79.07	15.00
98 TYR CA	-15.15	-18.84	79.78	15.00
98 TYR CB	-15.00	-20.35	79.61	15.00
98 TYR CG	-16.06	-21.14	80.35	15.00
98 TYR CD1	-17.39	-21.10	79.94	15.00
98 TYR CE1	-18.38	-21.80	80.65	15.00
98 TYR CD2	-15.74	-21.90	81.48	15.00
98 TYR CE2	-16.72	-22.60	82.18	15.00
98 TYR CZ	~18.03	-22.54	81.76	15.00
98 TYR OH	-19.00	-23.23	82.44	15.00
98 TYR C	-15.08	-18.52	81.26	15.00
98 TYR O	-14.04	-18.68	81.89	15.00
99 ASN N	-16.19	-18.01	81.79	15.00
99 ASN CA	-16.31	-17.67	83.19	15.00
99 ASN CB	-16.96	-16.29	83.38	15.00
99 ASN CG	-17.25	-15.98	84.84	15.00
99 ASN OD1	-16.67	-16.58	85.75	15.00
99 ASN ND2	-18.19	-15.06	85.07	15.00
99 ASN C	-17.20	-18.74	83.81	15.00
99 ASN O	-18.44	-18.61	83.82	15.00
100 PRO N	-16.59	-19.79	84.39	15.00
100 PRO CD	-15.15	-19.84	84.68	15.00
100 PRO CA	-17.29			15.00
100 PRO CB	-16.19		85.86	15.00
100 PRO CG		-20.47	86.03	15.00
100 PRO C		-20.55	85.87	15.00
100 PRO O		-21.29	85.92	15.00
101 THR N		-19.40	86.53	15.00
101 THR CA	-19.58	-19.00	87.37	15.00
101 THR CB	-19.22	-17.77	88.22	15.00
101 THR OG1	-18.93	-16.65	87.36	15.00
101 THR CG2		-18.06	89.08	15.00
101 THR C	-20.81	-18.70	86.52	15.00

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101 THR 0	-21.94	-19.07	86.88	15.00
102 GLY N	-20.59	-18.06	85.37	15.00
102 GLY CA	-21.68		84.47	15.00
102 GLY C	-22.27	-18.92	83.72	15.00
102 GLY O	-23.16	-18.73	82.87	15.00
103 LYS N	-21.80	-20.14	84.00	15.00
103 LYS CA	-22.32	-21.34	83.33	15.00
103 LYS CB	-21.70	-22.60	83.93	15.00
103 LYS CG	-22.23	-23.88	83.33	15.00
103 LYS CD	-21.78	-25.11	84.11	15.00
103 LYS CE	-22.56	-25.26	85.40	15.00
103 LYS NZ	-24.02	-25.47	85.12	15.00
103 LYS C	-23.83	-21.42	83.45	15.00
103 LYS O	-24.39	-21.26	84.53	15.00
104 ALA N	-24.50	-21.68	82.34	15.00
104 ALA CA	-25.96	-21.77	82.34	15.00
104 ALA CB	-26.56	-20.57	81.61	15.00
104 ALA C	-26.48	-23.07	81.73	15.00
104 ALA O	-27.69	-23.30	81.71	15.00
105 ALA N	-25.59	-23.92	81.24	15.00
105 ALA CA	-26.02	-25.18	80.65	15.00
105 ALA CB	-26.71	-24.94	79.31	15.00
105 ALA C	-24.89	-26.20	80.49	15.00
105 ALA O	-23.72	-25.84	80.50	15.00
106 LYS N	-25.28	-27.46	80.37	15.00
106 LYS CA	-24.36	-28.59	80.21	15.00
106 LYS CB	-24.28	-29.40	81.51	15.00
106 LYS CG	-23.38	-28.89	82.62	15.00
106 LYS CD	-23.65	-29.73	83.87	15.00
106 LYS CE	-22.48	-29.72	84.85	15.00
106 LYS NZ	-21.33	-30.54	84.36	15.00
106 LYS C	-24.99	-29.49	79.16	15.00
106 LYS O	-26.18	-29.32	78.83	15.00
107 CYS N	-24.23	-30.42	78.61	15.00
107 CYS CA	-24.77	-31.37	77.64	15.00
107 CYS CB	-24.82	-30.77	76.22	15.00
107 CYS SG	-23.38	-31.03	75.18	15.00
107 CYS C	-23.93	-32.64	77.72	15.00
107 CYS O	-22.76	-32.59	78.10	15.00
108 ARG N	-24.54	-33.78	77.45	15.00
108 ARG CA		-35.07	77.52	15.00
108 ARG CB	-24.60	-35.99	78.50	15.00
108 ARG CG	-26.12	-35.73	78.59	15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (Å<sup>2</sup>) for the cathepsin K complex with inhibitor 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide.

Residue Atom	x	Y	Z	В
1 ALA CB	-44.52	-37.54	64.26	15.00
1 ALA C	-46.72	-36.34	64.48	15.00
1 ALA O	-47.32	-36.96	63.59	15.00
1 ALA N	-46.03	-38.05	66.17	15.00
1 ALA CA	-45.55	-36.98	65.24	15.00
2 PRO N	-47.09	-35.10	64.86	15.00
2 PRO CD	-46.48	-34.27	65.92	15.00
2 PRO CA	-48.19	-34.39	64.20	15.00
2 PRO CB	-48.32	-33.13	65.04	15.00
2 PRO CG	-46.89	-32.89	65.50	15.00
2 PRO C	-47.85	-34.05	62.76	15.00
2 PRO O	-46.73	-34.29	62.29	15.00
3 ASP N	-48.84	-33.52	62.05	15.00
3 ASP CA	-48.64	-33.12	60.66	15.00
3 ASP CB	-49.97	-33.13	59.91	15.00
3 ASP CG	-50.31	-34.49	59.37	15.00
3 ASP OD1	-50.61	-34.58	58.16	15.00
3 ASP OD2	-50.25	-35.48	60.14	15.00
3 ASP C	-48.06	-31.73	60.63	15.00
3 ASP O	-47.45		59.63	15.00
4 SER N	-48.18	-31.03	61.75	15.00
4 SER CA	-47.72	-29.67	61.87	15.00
4 SER CB	-48.86	-28.74	61.51	15.00
4 SER OG	-48.48	-27.37	61.57	15.00
4 SER C	-47.29	-29.41	63.29	15.00
4 SER O	-47.89	-29.91	64.23	15.00
5 VAL N	-46.20	-28.66	63.43	15.00
5 VAL CA	-45.71	-28.29	64.73	15.00
5 VAL CB	-44.98	-29.44	65.47	15.00
5 VAL CG1	-43.59	-29.68	64.89	15.00
5 VAL CG2	-44.88	-29.11	66.96	15.00
5 VAL C	-44.81	-27.08	64.62	15.00
5 VAL O	-44.04		63.66	15.00
6 ASP N	-44.96	-26.20	65.59	15.00
6 ASP CA	-44.19	-24.98	65.66	15.00

6 ASP CB	-45.08	-23.81	<b>65 22</b>	15.00
6 ASP CG	-44.31	-22.52	65.22 65.06	15.00
6 ASP OD1	-43.13		65.49	15.00
6 ASP OD2	-44.88	-21.58	64.49	15,00
6 ASP C	-43.72	-24.81	67.10	15.00
6 ASP O	-44.50	-24.48	68.00	15.00
7 TYR N	-42.42		67.31	15.00
7 TYR CA	-41.83	-24.90	68.64	15.00
7 TYR CB	-40.43	-25.53	68.66	15.00
7 TYR CG	-40.49	-27.05	68.76	15.00
7 TYR CD1	-40.75	-27.66	69.98	15.00
7 TYR CE1	-40.88	-29.04	70.08	15.00
7 TYR CD2	-40.34	-27.85	67.63	15.00 15.00
7 TYR CE2	-40.47	-29.24	67.72	15.00
7 TYR CZ	-40.74	-29.83	68.95	15.00
7 TYR OH	-40.89	-31.20	69.06	15.00
7 TYR C	-41.80	-23.47	69.20	15.00
7 TYR O	-41.66	-23.28	70.42	15.00
8 ARG N	-41.93	-22.48	68.33	15.00
8 ARG CA	-41.95	-21.08	68.77	15.00
8 ARG CB	-42.06	-20.12	67.58	15.00
8 ARG CG	-40.92	-20.21	66.57	15.00
8 ARG CD	-41.19	-19.30	65.38	15.00
8 ARG NE	-42.23	-19.84	64.51	15.00
8 ARG CZ	-42.66	-19.26	63.38	15.00
8 ARG NH1	-42.13	-18.11	62.97	15.00
8 ARG NH2	-43.61	-19.84	62.66	15.00
8 ARG C	-43.20	-20.93	69.64	15.00
8 ARG O	-43.18	-20.25	70.68	15.00
9 LYS N	-44.28	-21.58	69.21	15.00
9 LYS CA	-45.54	-21.53	69.92	15.00
9 LYS CB	-46.66	-22.14	69.08	15.00
9 LYS CG	-47.11	-21.26	67.92	15.00
9 LYS CD		-22.07	66.95	15.00
9 LYS CE	-48.74	-21.20	65.98	15.00
9 LYS NZ	-49.84	-20.49	66.67	15.00
9 LYS C	-45.45	-22.22	71.27	15.00
9 LYS O	-46.19	-21.88	72.19	15.00
10 LYS N	-44.53	-23.16	71.39	15.00
10 LYS CA	-44.30	-23.91	72.63	15.00
10 LYS CB	-43.82	-25.33	72.30	15.00
10 LYS CG	-44.90	-26.25	71.75	15.00
10 LYS CD	-44.35	-27.64	71.47	15.00

10	LYS	CE	-45.48	-28.63	71.20	15.00
10	LYS	NZ	-44.99	-30.01	70.87	15.00
10	LYS	С	-43.28	-23.22	73.53	15.00
10	LYS	0	-42.94	-23.73	74.60	15.00
11	GLY	N	-42.75	-22.09	73.09	15.00
11	GLY	CA	-41.77	-21.37	73.88	15.00
11	GLY	С	-40.41	-22.04	73.97	15.00
11	GLY	0	-39.71	-21.91	74.97	15.00
12	TYR	N	-40.02	-22.75	72.92	15.00
12	TYR	CA	-38.73	-23.41	72.89	15.00
12	TYR	CB	-38.86	-24.81	72.29	15.00
12	TYR	CG	-39.47	-25.86	73.18	15.00
12	TYR	CD1	-40.56	-25.59	73.99	15.00
12	TYR	CE1	-41.12	-26.56	74.81	15.00
12	TYR	CD2	-38.94	-27.15	73.21	15.00
12	TYR	CE2	-39.49	-28.14	74.02	15.00
12	TYR	CZ	-40.58	-27.84	74.82	15.00
12	TYR	OH	-41.10	-28.81	75.64	15.00
12	TYR	C	-37.73	-22.62	72.04	15.00
12	TYR	0	-36.65	-23.14	71.72	15.00
13	VAL	N	-38.08	-21.39	71.66	15.00
13	VAL	CA	-37.21	-20.59	70.80	15.00
13	VAL	CB	-37.82	-20.46	69.40	15.00
13	VAL	CG1	-36.75	-20.07	68.38	15.00
13	VAL	CG2	-38.52	-21.73	69.02	15.00
13	VAL	С	-36.93	-19.17	71.30	15.00
13	VAL	0	-37.86	-18.42	71.61	15.00
14	THR	N	-35.66	-18.79	71.34	15.00
14	THR	CA	-35.29	-17.45	71.78	15.00
14	THR	CB	-33.84	-17.40	72.32	15.00
14	THR	OG1	-32.91	-17.76	71.28	15.00
14	THR	CG2	-33.67	-18.33	73.52	15.00
14	THR	C	-35.46	-16.52	70.59	15.00
14	THR	0	-35.55	-16.96	69.46	15.00
15	PRO	N	-35.49	-15.20	70.84	15.00
15	PRO	CD	-35.39	-14.48	72.12	15.00
15	PRO	CA	-35.65	-14.27	69.72	15.00
15	PRO	CB	-35.71	-12.90	70.42	15.00
15	PRO	CG	-34.93	-13.12	71.67	15.00
15	PRO	С	-34.54	-14.35	68.69	15.00
15	PRO	0	-33.45	-14.89	68.95	15.00
16	VAL	N	-34.85	-13.85	67.50	15.00
16	VAL	CA	-33.94	-13.84	66.37	15.00

16 VAL CB	-34.68	-13.41	65.08	15.00
16 VAL CG1	-33.72	-13.35	63.90	15.00
16 VAL CG2	-35.82	-14.39	64.78	15.00
16 VAL C	-32.71	-12.94	66.61	15.00
16 VAL 0	-32.84	-11.79	67.04	15.00
17 LYS N	-31.54	-13.50	66.33	15.00
17 LYS CA	-30.27	-12.80	66.48	15.00
17 LYS CB	-29.26	-13.70	67.22	15.00
17 LYS CG	-29.85	-14.45	68.41	15.00
17 LYS CD	-30.15	-13.52	69.56	15.00
17 LYS CE	-31.10	-14.14	70.58	15.00
17 LYS NZ	-30.73	-15.52	70.94	15.00
17 LYS C	-29.70	-12.43	65.09	15.00
17 LYS 0	-30.25	-12.81	64.05	15.00
18 ASN N	-28.59	-11.71	65.10	15.00
18 ASN CA	-27.93	-11.24	63.89	15.00
18 ASN CB	-28.25	-9.75	63.69	15.00
18 ASN CG	-27.74	-9.21	62.37	15.00
18 ASN OD1	-27.34	-9.95	61.49	15.00
18 ASN ND2	-27.75	-7.90	62.24	15.00
18 ASN C	-26.41	-11.47	63.99	15.00
18 ASN 0	-25.70	-10.74	64.68	15.00
19 GLN N	-25.92	-12.47	63.27	15.00
19 GLN CA	-24.50	-12.81	63.27	15.00
19 GLN CB	-24.23	-14.05	62.39	15.00
19 GLN CG	-24.59	-13.91	60.91	15.00
19 GLN CD	-24.31	-15.17	60.10	15.00
19 GLN OE1	-25.15	-16.06	60.00	15.00
19 GLN NE2	-23.13	-15.23	59.50	15.00
19 GLN C	-23.59	-11.65	62.84	15.00
19 GLN O	-22.45	-11.56	63.30	15.00
20 GLY N	-24.09	-10.77	61.99	15.00
20 GLY CA	-23.28	-9.65	61.53	15.00
20 GLY C	-22.31	-10.07	60.45	15.00
20 GLY O	-22.59	-10.98	59.67	15.00
21 GLN N .	-21.15	-9.43	60.41 .	15.00
21 GLN CA	-20.14	-9.75	59.41	15.00
21 GLN CB	-19.40	-8.48	58.96	15.00
21 GLN CG	-20.22	-7.59	58.06	15.00
21 GLN CD	-20.48	-8.23	56.69	15.00
21 GLN 0E1	-19.88	-9.26	56.33	15.00
21 GLN NE2	-21.37	-7.60	55.90	15.00
21 GLN C	-19.15	-10.78	59.98	15.00

21	GLN	0	-17.95	-10.51	60.13	15.00
22	CYS	N	-19.68	-11.95	60.30	15.00
22	CYS	CA	-18.90	-13.04	60.86	15.00
22	CYS	С	-19.59	-14.32	60.46	15.00
22	CYS	0	-20.82	-14.40	60.50	15.00
22	CYS	CB	-18.83	-12.90	62.40	15.00
22	CYS	SG	-18.13	-14.32	63.33	15.00
23	GLY	N	-18.81	-15.30	60.03	15.00
23	GLY	CA	-19.36	-16.58	59.63	15.00
23	GLY	С	-19.61	-17.48	60.83	15.00
23	GLY	0	-19.23	-18.65	60.84	15.00
24	SER	N	-20.32	-16.94	61.82	15.00
24	SER	CA	-20.63	-17.67	63.03	15.00
24	SER	CB	-20.58	-16.71	64.22	15.00
24	SER	OG	-21.38	-15.58	64.00	15.00
24	SER	C	-22.00	-18.34	62.98	15.00
24	SER	0	-22.52	-18.78	64.01	15.00
	CYS		-22.59	-18.45	61.79	15.00
	CYS		-23.90	-19.08	61.65	15.00
	CYS		-24.31	-19.15	60.17	15.00
	CYS		-23.12	-20.00	59.06	15.00
25		С	-23.95	-20.47	62.29	15.00
25			-24.95	-20.85	62.89	15.00
25			-28.28	-9.31	55.94	15.00
25			-28.07	-9.03	57.30	15.00
25			-27.11	-9.78	58.03	15.00
25			-26.37	-10.78	57.40	15.00
25			-26.59	-11.05	56.05	15.00
25			-27.54	-10.32	55.31	15.00
25			-25.31	-11.54	58.16	15.00
25			-24.19	-11.68	57.24	15.00
25		C9	-23.29	-12.79	57.20	15.00 15.00
25		010	-22.50	-12.99	58.13	15.00
	INH		-22.45		55.88	
		C12	-21.05	-14.47	56.48	15.00 15.00
25		C13	-20.11	-13.38	55.92	15.00
		C14	-19.15	-12.91	57.01	15.00
25		C15	-20.83	-12.17	55.23 56.34	15.00
25		C16	-23.00	-16.06	56.75	15.00
25		017	-24.16	-16.15	56.30	15.00
25		N18	-22.19	-17.17	56.74	15.00
25		N19	-22.62	-18.53		15.00
25	INH	N20	-23.34	-13.55	56.10	13.00

25 INH C21		-18.85	58.14	15.00
25 INH 022	-22.10		58.75	15.00
25 INH C23			59.77	15.00
25 INH C24	-		60.62	15.00
25 INH C25	-14.61	-	60.17	15.00
25 INH C26	-14.52	-25.29	58.88	15.00
25 INH C27	-13.54	-25.80	58.03	15.00
25 INH C28	-12.67	-26.81	58.47	15.00
25 INH C29	-15.45	-24.21	58.40	15.00
25 INH 030	-16.52	-24.58	57.49	15.00
25 INH C31	-17.56	-23.66	57.05	15.00
25 INH 032	-17.32	-22.74	56.27	15.00
25 INH C33	-19.95		57.18	15.00
25 INH C34	-21.23	-23.90	57.35	15.00
25 INH C35	-21.11	-25.25	58.03	15.00
25 INH C36	-22.32	-25.59	58.89	15.00
25 INH C37	-20.84		56.99	15.00
25 INH C38	-20.07	-21.83	58.03	15.00
25 INH 039	-19.74	-21.86	59.22	15.00
25 INH N40	-20.56	-20.70	57.43	15.00
25 INH N41	-20.70	-19.44	58.21	15.00
25 INH N42	-18.78	-23.90	57.54	15.00
26 TRP N	-22.83	-21.19	62.21	15.00
26 TRP CA	-22.70	-22.53	62.79	15.00
26 TRP CB	-21.33	-23.13	62.41	15.00
26 TRP CG	-20.12	-22.32	62.88	15.00
26 TRP CD2	-19.43	-22.44	64.14	15.00
26 TRP CE2	-18.42	-21.46	64.16	15.00
26 TRP CE3	-19.58	-23.27	65.26	15.00
26 TRP CD1	-19.50	-21.31	62.20	15.00
26 TRP NE1	-18.48	-20.79	62.96	15.00
26 TRP CZ2	-17.56	-21.29	65.25	15.00
26 TRP CZ3	-18.73			15.00
26 TRP CH2	-17.73	-22.11	66.33	15.00
26 TRP C	-22.87	-22.47	64.31	15.00
26 TRP O	-23.46	-23.3 <i>6</i>	64.91	15.00
27 ALA N	-22.37	-21.39	64.90	15.00
27 ALA CA	-22.43	-21.17	66.34	15.00
27 ALA CB	-21.53	-20.00	66.72	15.00
27 ALA C	-23.87	-20.90	66.77	15.00
27 ALA O	-24.34	-21.42	67.78	15.00
28 PHE N	-24.55	-20.06	65.99	15.00
28 PHE CA	-25.94	-19.71	56.23	15.00
			<del>-</del>	<del></del>

28	PHE	CB	-26.38	-18.58	65.29	15.00
28	PHE	CG	-25.85	-17.23	65.70	15.00
28	PHE	CD1	-24.59	-16.81	65.30	15.00
28	PHE	CD2	-26.60	-16.39	66.52	15.00
28	PHE	CE1	-24.08	-15.57	65.70	15.00
28	PHE	CE2	-26.09	-15.15	66.93	15.00
28	PHE	CZ	-24.83	-14.74	66.52	15.00
28	PHE	С	-26.86	-20.93	66.07	15.00
28	PHE	0	-27.82	-21.10	66.82	15.00
29	SER	N	-26.54	-21.81	65.13	15.00
29	SER	CA	-27.33	-23.01	64.93	15.00
29	SER	CB	-26.85	-23.73	63.66	15.00
29	SER	OG	-27.55	-24.93	63.47	15.00
29	SER	С	-27.19	-23.93	66.16	15.00
29	SER	0	-28.19	-24.39	66.71	15.00
30	SER	N	-25.95	-24.16	66.59	15.00
30	SER	CA	-25.63	-25.00	67.75	15.00
30	SER	CB	-24.12	-24.97	68.04	15.00
30	SER	OG	-23.34	-25.28	66.91	15.00
30	SER	С	-26.36	-24.51	68.98	15.00
30	SER	0	-27.02	-25.26	69.69	15.00
31	VAL	N	-26.16	-23.23	69.26	15.00
31	VAL	CA	-26.76	-22.55	70.39	15.00
31	VAL	CB	-26.31	-21.07	70.38	15.00
31	VAL	CG1	-27.42	-20.13	70.84	15.00
31	VAL	CG2	-25.08	-20.92	71.26	15.00
31	VAL	С	-28.28	-22.70	70.37	15.00
31	VAL	0	-28.87	-23.11	71.37	15.00
32	GLY	N	-28.89	-22.44	69.22	15.00
32	GLY	CA	-30.32	-22.56	69.08	15.00
32	GLY	С	-30.79	-23.96	69.45	15.00
32	GLY	0	-31.80	-24.12	70.14	15.00
33	ALA	N	-30.03	-24.96	69.01	15.00
33	ALA	CA	-30.35	-26.35	69.30	15.00
33.	ALA	CB	-29.44	-27.29	68.48	15.00
33	ALA	С	-30.24	-26.64	70.80	15.00
33	ALA	0	-31.11	-27.29	71.38	15.00
34	LEU	N	-29.17	-26.15	71.42	15.00
34	LEU	CA	-28.96	-26.37	72.84	15.00
34	LEU	CB	-27.61	-25.92	73.28	15.00
34	LEU	CG	-26.39	-26.66	72.87	15.00
34	LEU	CD1	-25.12	-25.82		15.00
34	LEU	CD2	-26.27	-27.87	73.78	15.00

37 GLN NE2 -34.16 -30.78 71.87 15.00 37 GLN C -33.10 -28.48 76.17 15.00 37 GLN O -33.70 -29.09 77.06 15.00 38 LEU N -32.38 -27.39 76.40 15.00						
34 LEU O	34	LEU C	-30.09	-25.73	73.66	15.00
35 GLU N	34	LEU O	-30.62			
35 GLU CA	35	GLU N	-30.48	-24.53		
35 GLU CB	35	GLU CA	-31.55	-23.78		
35 GLU CG	35	GLU CB	-31.70			
35 GLU CD	35	GLU CG	-30.49			
35 GLU OE1	35	GLU CD	-30.59	-20.25		
35 GLU OE2	35	GLU OE1	-31.42	-20.17		
35 GLU C	35	GLU OE2	-29.81	-19.32		
35 GLU O	35	GLU C	-32.89			
36 GLY N	35	GLU O		-24.41		
36 GLY CA       -34.41       -25.97       72.72       15.00         36 GLY C       -34.42       -27.05       73.78       15.00         36 GLY O       -35.32       -27.13       74.61       15.00         37 GLN N       -33.35       -27.86       73.77       15.00         37 GLN CA       -33.18       -28.95       74.71       15.00         37 GLN CB       -31.95       -29.77       74.32       15.00         37 GLN CG       -32.01       -30.34       72.91       15.00         37 GLN CD       -33.22       -31.25       72.69       15.00         37 GLN OE1       -33.31       -32.35       73.26       15.00         37 GLN NE2       -34.16       -30.78       71.87       15.00         37 GLN C       -33.10       -28.48       76.17       15.00         38 LEU N       -32.38       -27.39       76.40       15.00         38 LEU CA       -32.27       -26.87       77.75       15.00         38 LEU CB       -31.39       -25.63       77.79       15.00         38 LEU CD       -30.49       -25.87       80.11       15.00         38 LEU CD       -33.65       -26.51 <td< td=""><td>36</td><td>GLY N</td><td>-33.16</td><td>-25.25</td><td></td><td></td></td<>	36	GLY N	-33.16	-25.25		
36 GLY C	36	GLY CA				
36 GLY O	36	GLY C	-34.42	-27.05		
37 GLN N	36	GLY O	-35.32	-27.13		
37 GLN CA	37	GLN N	-33.35	-27.86		
37 GLN CB	37	GLN CA	-33.18	-28.95		
37 GLN CG	37	GLN CB	-31.95	-29.77		
37 GLN CD	37	GLN CG	-32.01	-30.34	72.91	
37 GLN OE1	37	GLN CD	-33.22	-31.25		
37 GLN NE2	37	GLN OE1	-33.31	-32.35		15.00
37 GLN C	37	GLN NE2	-34.16	-30.78	71.87	
37 GLN O	37	GLN C	-33.10	-28.48	76.17	
38 LEU N	37	GLN O	-33.70	-29.09	77.06	15.00
38 LEU CB	38	LEU N	-32.38	-27.39	76.40	15.00
38 LEU CG	38	LEU CA	-32.27	-26.87	77.75	15.00
38 LEU CD1 -30.49 -25.87 80.11 15.00 38 LEU CD2 -30.79 -23.61 79.08 15.00 38 LEU C -33.65 -26.51 78.26 15.00 38 LEU C -33.97 -26.73 79.43 15.00 39 LYS N -34.45 -25.90 77.38 15.00 39 LYS CA -35.81 -25.51 77.72 15.00 39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	38	LEU CB	-31.39	-25.63	77.79	15.00
38 LEU CD2 -30.79 -23.61 79.08 15.00 38 LEU C -33.65 -26.51 78.26 15.00 38 LEU O -33.97 -26.73 79.43 15.00 39 LYS N -34.45 -25.90 77.38 15.00 39 LYS CA -35.81 -25.51 77.72 15.00 39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	38	LEU CG	-31.34	-25.01	79.18	15.00
38 LEU C -33.65 -26.51 78.26 15.00 38 LEU O -33.97 -26.73 79.43 15.00 39 LYS N -34.45 -25.90 77.38 15.00 39 LYS CA -35.81 -25.51 77.72 15.00 39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS CE -39.79 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	38	LEU CD1	-30.49	-25.87	80.11	15.00
38 LEU O -33.97 -26.73 79.43 15.00 39 LYS N -34.45 -25.90 77.38 15.00 39 LYS CA -35.81 -25.51 77.72 15.00 39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	38	LEU CD2	-30.79	-23.61	79.08	15.00
39 LYS N -34.45 -25.90 77.38 15.00 39 LYS CA -35.81 -25.51 77.72 15.00 39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS C -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	38	LEU C	-33.65	-26.51	78.26	15.00
39 LYS CA -35.81 -25.51 77.72 15.00 39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS C -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	38	LEU O	-33.97	-26.73	79.43	15.00
39 LYS CB -36.42 -24.72 76.55 15.00 39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00			-34.45	-25.90	77.38	15.00
39 LYS CG -37.63 -23.88 76.91 15.00 39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	3.9	LYS CA	-35.81	-25.51	77.72	15.00
39 LYS CD -38.88 -24.71 77.08 15.00 39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	39	LYS CB	-36.42	-24.72	76.55	15.00
39 LYS CE -39.79 -24.14 78.15 15.00 39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	39	LYS CG	-37.63	-23.88	76.91	15.00
39 LYS NZ -39.99 -22.66 78.01 15.00 39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	39	LYS CD	-38.88	-24.71	77.08	15.00
39 LYS C -36.61 -26.78 78.00 15.00 39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	39	LYS CE	-39.79	-24.14	78.15	15.00
39 LYS O -37.28 -26.89 79.04 15.00 40 LYS N -36.48 -27.75 77.11 15.00	39	LYS NZ	-39.99	-22.66	78.01	15.00
40 LYS N -36.48 -27.75 77.11 15.00	39	LYS C	-36.61	-26.78	78.00	15.00
40 LYS N -36.48 -27.75 77.11 15.00	39	LYS O	-37.28	-26.89	79.04	15.00
40 LYS CA -37.19 -29.01 77.23 15.00	40	LYS N	-36.48	-27.75	77.11	15.00
	40	LYS CA	-37.19	-29.01	77.23	15.00

40	LYS	CB	-36.93	-29.90	76.01	15.00
40	LYS	CG	-37.84	-31.13	75.91	15.00
40	LYS	CD	-37.59	-31.87	74.60	15.00
40	LYS	CE	-37.06	-33.28	74.85	15.00
40	LYS	NZ	-36.36	-33.87	73.66	15.00
40	LYS	C	-36.82	-29.76	78.51	15.00
40	LYS	0	-37.69	-30.36	79.16	15.00
41	LYS	N	-35.55	-29.66	78.92	15.00
41	LYS	CA	-35.08	-30.37	80.10	15.00
41	LYS	CB	-33.60	-30.71	79.95	15.00
41	LYS	CG	-33.12	-31.74	80.95	15.00
41	LYS	CD	-31.66	-32.10	80.76	15.00
41	LYS	CE	-31.18	-33.02	81.87	15.00
41	LYS	NZ	-31.86	-34.35	81.80	15.00
41	LYS	С	-35.32	-29.68	81.45	15.00
41	LYS	0	-35.76	-30.32	82.40	15.00
42	THR	N	-35.04	-28.38	81.51	15.00
42	THR	CA	-35.19	-27.62	82.75	15.00
42	THR	CB	-34.00	-26.66	82.95	15.00
42	THR	OG1	-34.10	-25.58	82.01	15.00
42	THR	CG2	-32.68	-27.39	82.71	15.00
42	THR	С	-36.46	-26.78	82.86	15.00
42	THR	0	-36.86	-26.41	83.96	15.00
43	GLY	N	-37.04	-26.42	81.73	15.00
43	GLY	CA	-38.24	-25.60	81.75	15.00
43	GLY	C	-37.95	-24.12	81.59	15.00
43	GLY	0	-38.88	-23.31	81.62	15.00
44	LYS	N	-36.67	-23.75	81.48	15.00
44	LYS	CA	-36.28	-22.35	81.31	15.00
44	LYS	CB	-35.14	-21.96	82.26	15.00
44	LYS		-35.58	-21.64	83.67	15.00
44	LYS		-35.80	-22.90	84.47	15.00
	LYS		-34.48	-23.49	84.91	15.00
	LYS		-33.81	-22.59	85.90	15.00
	LYS	С	-35.82	-22.07	79.87	15.00
44	LYS	0	-35.33	-22.95	79.19	15.00
45	LEU		-35.97	-20.82	79.44	15.00
45	LEU	CA	-35.56	-20.40	78.11	15.00
45	LEU		-36.79	-20.00	77.26	15.00
45	ľŒŪ		-36.54	-19.61	75.80	15.00
	LEU		-36.32	-20.88	74.98	15.00
	LEU		-37.71	-18.81	75.22	15.00
45	LEU	C	-34.65	-19.18	78.26	15.00

45 LE	U O	-35.09	-18.11	78.69	15.00
46 LE	UN	-33.36	-19.38	78.00	15.00
46 LE	U CA	-32.41	-18.30	78.06	15.00
46 LE	U CB	-31.64	-18.25	79.40	15.00
46 LE	U CG	-30.80	-19.34	80.08	15.00
46 LE	U CD1	-31.61	-20.01	81.15	15.00
46 LET	U CD2	-30.27	-20.35	79.09	15.00
46 LET	J C	-31.46	-18.38	76.86	15.00
46 LET	O	-31.39	-19.40	76.17	15.00
47 ASI	N	-30.79	-17.27	76.59	15.00
47 ASN	1 CA	-29.86	-17.18	75.47	15.00
47 ASN	1 CB	-29.74	-15.73	75.04	15.00
47 ASN	1 CG	-31.07	-15.14	74.66	15.00
47 ASN	N OD1	-31.74	-15.64	73.75	15.00
47 ASN	ND2	-31.50	-14.13	75.39	15.00
47 ASN	1 C	-28.51	-17.75	75.81	15.00
47 ASN	10	-27.91	-17.37	76.81	15.00
	N	-28.07	-18.74	75.05	15.00
	CA	-26.77	-19.34	75.27	15.00
	CB	-26.81	-20.84	74.95	15.00
	CG	-27.74	-21.60	75.90	15.00
	CD1	-27.67	-23.09	75.67	15.00
	CD2	-27.37	-21.30	77.33	15.00
48 LEU		-25.77	-18.57	74.42	15.00
48 LEU		-26.17	-17.88	73.47	15.00
49 SER		-24.50	-18.67	74.76	15.00
	CA	-23.45	-17.92	74.09	15.00
	CB	-22.32	-17.62	75.08	15.00
	OG	-21.28	-16.87	74.48	15.00
49 SER		-22.83	-18.44	72.80	15.00
49 SER		-22.08	-19.40	72.82	15.00
50 PRO		-23.10	-17.78	71.67	15.00
	CD	-24.13		71.42	15.00
50 PRO			-18.23	70.41	15.00
	CB		-17.43	69.35	15.00
50 PRO		-24.58	-17.10	70.04	15.00
50 PRO		-21.01	-17.85	70.42	15.00
50 PRO		-20.16	-18.58	69.92	15.00
51 GLN		-20.70	-16.70	71.03	15.00
51 GLN		-19.34	-15.20	71.12	15.00
51 GLN		-19.32	-14.84	71.83	15.00
51 GLN		-18.00	-14.10	71.73	15.00
51 GLN	CD	-1772	-13.59	70.34	15.00

51	GLN	OE1	-18.55	-12.90	69.73	15.00
51	GLN	NE2	-16.55	-13.90	69.82	15.00
51	GLN	С	-18.45	-17.21	71.86	15.00
51	GLN	0	-17.28	-17.39	71.50	15.00
52	ASN	N	-19.01	-17.87	72.87	15.00
52	ASN	CA	-18.28	-18.88	73.62	15.00
52	ASN	CB	-19.24	-19.57	74.60	15.00
52	ASN	CG	-18.54	-20.47	75.63	15.00
52	ASN	OD1	-19.19	~20.95	76.57	15.00
52	ASN	ND2	-17.24	-20.70	75.48	15.00
52	ASN	С	-17.74	-19.88	72.59	15.00
52	ASN	0	-16.55	-20.22	72.60	15.00
53	LEU	N	-18.60	-20.31	71.68	15.00
53	LEU	CA	-18.22	-21.26	70.64	15.00
53	LEU	CB	-19.47	-21.70	69.86	15.00
53	LEU	CG	-20.37	-22.81	70.42	15.00
53	LEU	CD1	-20.05	-23.13	71.87	15.00
53	LEU	CD2	-21.83	-22.41	70.22	15.00
53	LEU	С	-17.18	-20.69	69.68	15.00
53	LEU	0	-16.17	-21.33	69.40	15.00
54	VAL	N	-17.44	-19.48	69.18	15.00
54	VAL	CA	-16.55	-18.82	68.23	15.00
54	VAL	CB	-17.03	-17.36	67.92	15.00
54	VAL	CG1	-15.96	-16.58	67.19	15.00
54	VAL	CG2	-18.28	-17.40	67.05	15.00
54	VAL	C	-15.10	-18.79	68.72	15.00
54	VAL	0	-14.19	-19.17	67.98	15.00
55	ASP	N	-14.90	-18.37	69.96	15.00
55	ASP	CA	-13.56	-18.26	70.51	15.00
55			-13.56	-17.25	71.67	15.00
55	ASP	CG	-14.04	-15.87	71.27	15.00
	ASP		-14.02	-15.54	70.06	15.00
	ASP		-14.43	-15.10	72.18	15.00
	ASP		-12.93	-19.55	71.04	15.00
	ASP	0	-11.72	-19.70	71.00	15.00
56	CYS	N	<b>-13.76</b>	-20.47	71.50	15.00
56	CYS	CA	-13.26	-21.69	72.14	15.00
56	CYS	C	-13.18	-23.02	71.37	15.00
	CYS	0	-12.34	-23.86	71.72	15.00
56	CYS	CB	-14.03	-21.89	73.45	15.00
56	CYS	SG	-14.05	-20.41	74.53	15.00
57	VAL	N	-14.03	-23.22	70.37	15.00
57	VAL	CA	-14.00	-24.47	69.62	15.00

	VAL		-15.34	-24.76	68.86	15.00
57		CG1	-15.38	-26.20	68.38	15.00
57	VAL	CG2	-16.52	-24.51	69.77	15.00
57	VAL	С	-12.80	-24.45	68.66	15.00
57	VAL	0	-12.85	-23.83	67.60	15.00
58	SER	N	-11.72	-25.10	69.08	15.00
58	SER	CA	-10.48	-25.16	68.32	15.00
58	SER	CB	-9.34	-25.68	69.21	15.00
58	SER	OG	-9.70	-26.91	69.82	15.00
58	SER	С	-10.53	-25.97	67.03	15.00
58	SER	0	-9.67	-25.81	66.17	15.00
59	GLU	N	-11.49	-26.88	66.92	15.00
59	GLU	CA	-11.62	-27.68	65.70	15.00
59	GLU	CB	-12.33	-29.00	65.99	15.00
	GLU		-11.57	-29.96	66.92	15.00
59	GLU	CD	-11.66	-29.59	68.40	15.00
	GLU		-10.69	-29.88	69.12	15.00
59	GLU	OE2	-12.69	-29.03	68.85	15.00
59	GLU	C	-12.37	-26.89	64.62	15.00
	GLU		-12.48	-27.34	63.49	15.00
	ASN	N	-12.91	-25.73	64.98	15.00
60	ASN	CA	-13.65	-24.87	64.06	15.00
	ASN	CB	-14.99	-24.44	64.66	15.00
	ASN	CG	-16.08	-25.51	64.51	15.00
	ASN		-17.16	-25.38	65.08	15.00
	ASN		-15.79	-26.55	63.74	15.00
	ASN		-12.81	-23.65	63.73	15.00
	asn		-11.76	-23.44	64.35	15.00
	ASP		-13.28	-22.85	62.78	15.00
	ASP		-12.53	-21.67	62.38	15.00
	ASP		-12.22	-21.74	60.88	15.00
	ASP		-11.54	-23.04	60.50	15.00
	ASP		-12.02	-23.72	59.56	15.00
	ASP		-10.54		61.16	15.00
	ASP		-13.13		62.74	15.00
	ASP		-12.89	-19.32	62.04	15.00
	GLY		-13.88	-20.27	63.83	15.00
	GLY		-14.48	-19.03	64.30	15.00
	GLY		-15.34	-18.30	63.30	15.00
	GLY		-16.37	-18.81	62.87	15.00
	CYS		-14.96		62.96	
	CYS			-16.28	61.99	
63	CYS	С	-15.49	-16.79	60.56	15.00

63	CYS	0	-16.19	-16.39	59.63	15.00
63	CYS	CB	-15.37	-14.79	62.10	15.00
63	CYS	SG	-16.14	-13.94	63.53	15.00
64	GLY	N	-14.55	-17.71	60.41	15.00
64	GLY	CA	-14.27	-18.27	59.10	15.00
64	GLY	С	-15.11	-19.50	58.79	15.00
64	GLY	0	-14.95	-20.10	57.73	15.00
65	GLY	N	-15.94	-19.92	59.74	15.00
65	GLY	CA	-16.78	-21.08	59.52	15.00
65	GLY	С	-16.54	-22.20	60.51	15.00
65	GLY	0	-15.54	-22.20	61.24	15.00
66	GLY	N	-17.44	-23.18	60.52	15.00
66	GLY	CA	-17.29	-24.31	61.41	15.00
66	GLY	С	-18.41	-25.32	61.27	15.00
66	GLY	0	-19.25	-25.19	60.37	15.00
67	TYR	N	-18.41	-26.34	62.13	15.00
67	TYR	CA	-19.42	-27.39	62.15	15.00
67	TYR	CB	-18.78	-28.74	61.83	15.00
67	TYR	CG	-18.30	-28.84	60.41	15.00
67	TYR	CD1	-19.02	-29.55	59.47	15.00
67	TYR	CE1	-18.62	-29.59	58.13	15.00
67	TYR	CD2	-17.15	-28.16	60.00	15.00
67	TYR	CE2	-16.74	-28.19	58.67	15.00
67	TYR	CZ	-17.48	-28.90	57.74	15.00
67	TYR	OH	-17.12	-28.88	56.42	15.00
67	TYR	С	-20.12	-27.46	63.51	15.00
67	TYR	0	-19.48	-27.31	64.54	15.00
68	MET	N	-21.43	-27.73	63.50	15.00
68	MET	CA	-22.22	-27.81	64.73	15.00
68	MET	CB	-23.72	-27.83	64.41	15.00
68	MET	CG	-24.26	-26.60	63.63	15.00
68	MET	SD	-24.05	-26.62	61.79	15.00
68	MET		-25.42	-27.66	61.31	15.00
68	MET	C	-21.84	-29.02	65.61	15.00
68	MET	0	-21.89	-28.95	66.84	15.00
69	THR	N	-21.48	-30.12	64.98	15.00
69	THR	CA	-21.08	-31.33	65.70	15.00
69	THR	CB	-20.80	-32.51	64.73	15.00
69	THR	OG1	-20.08	-32.02	63.58	15.00
69	THR	CG2	-22.12	-33.15	64.27	15.00
69	THR	С	-19.82	-31.05	66.53	15.00
69	THR	0	-19.67	-31.55	67.66	15.00
70	ASN	N	-18.94	-30 21	65.99	15.00

70 ASN CA	-17.72	-29.85	66 60	15 00
70 ASN CB	-16.76		66.69	
70 ASN CG	-15.89	-29.97	65.79 64.95	15.00
70 ASN OD1	-15.35	-29.56	63.93	15.00
70 ASN ND2	-15.72	-31.21	65.39	15.00
70 ASN C	-18.08	-29.02	67.91	15.00
70 ASN O	-17.57	~29.25	69.00	15.00
71 ALA N	-19.02	-28.09	67.71	15.00 15.00
71 ALA CA	-19.50	-27.22	68.77	15.00
71 ALA CB	-20.49	-26.20	68.21	15.00
71 ALA C	-20.14	-28.02	69.90	15.00
71 ALA O	-19.88	-27.76	71.07	15.00
72 PHE N	-20.90	-29.05	69.55	15.00
72 PHE CA	-21.56	-29.89	70.55	15.00
72 PHE CB	-22.55	-30.86	69.89	15.00
72 PHE CG	-23.72	-30.18	69.24	15.00
72 PHE CD1	-24.30		69.82	15.00
72 PHE CD2	-24.23	-30.65	68.04	15.00
72 PHE CE1	-25.38	-28.41	69.21	15.00
72 PHE CE2	-25.31	-30.01		15.00
72 PHE CZ	-25.89	-28.89	68.01	15.00
72 PHE C	-20.53	-30.68	71.37	15.00
72 PHE 0	-20.66	-30.76	72.60	15.00
73 GLN N	-19.55	-31.27	70.69	15.00
73 GLN CA	-18.50	-32.05	71.34	15.00
73 GLN CB	-17.52	-32.66	70.34	15.00
73 GLN CG	-18.09	-33.80	69.51	15.00
73 GLN CD	-17.03	-34.63	68.79	15.00
73 GLN 0E1	-17.20	-35.01	67.62	15.00
73 GLN NE2	-15.95	-34.94	69.49	15.00
73 GLN C	-17.75	-31.16	72.30	15.00
73 GLN 0	-17.35	-31.61	73.38	15.00
74 TYR N	-17.55	-29.89	71.92	15.00
74 TYR CA	-16.84	-28.94	72.75	15.00
74 TYR CB	-16.75	-27.57	72.08	15.00
74 TYR CG	-16.46	-26.43	73.03	15.00
74 TYR CD1	-15.18	-26.26	73.58	15.00
74 TYR CE1	-14.91	-25.23	74.50	15.00
74 TYR CD2	-17.46	-25.54	73.42	15.00
74 TYR CE2	-17.21	-24.51	74.34	15.00
74 TYR CZ	-15.93	-24.36	74.88	15.00
74 TYR OH	-15.69	-23.36	75.78	15.00
74 TYR C	-17.58	-28.81	74.06	15.00

74	TYR	0	-17.02	-29.09	75.12	15.00
7.5	VAL	N	-18.84	-28.39	73.99	15.00
.75	VAL	CA	-19.67	-28.21	75.17	15.00
75	VAL	CB	-21.14	-27.89	74.77	15.00
75	VAL	CG1	-22.00	-27.77	76.01	15.00
75	VAL	CG2	-21.20	-26.59	73.96	15.00
75	VAL	С	-19.61	-29.43	76.09	15.00
75	VAL	0	-19.55	-29.27	77.31	15.00
76	GLN	N	-19.56	-30.63	75.51	15.00
76	GLN	CA	-19.48	-31.85	76.29	15.00
76	GLN	CB	-19.68	-33.09	75.42	15.00
76	GLN	CG	-19.53	-34.39	76.19	15.00
76	GLN	CD	-19.73	-35.63	75.35	15.00
76	GLN	OE1	-19.39	-35.66	74.17	15.00
76	GLN	NE2	-20.27	-36.67	75.97	15.00
76	GLN	C	-18.15	-31.96	77.03	15.00
76	GLN	0	-18.14	-32.20	78.23	15.00
77	LYS	N	-17.05	-31.81	76.30	15.00
77	LYS	CA	-15.70	-31.90	76.87	15.00
77	LYS	CB	-14.63	-31.71	75.79	15.00
77	LYS	CG	-14.73	-32.63	74.61	15.00
77	LYS	CD	-14.44	-34.07	74.97	15.00
77	LYS	CE	-14.49	-34.96	73.73	15.00
77	LYS	NZ	-13.56	-34.52	72.63	15.00
77	LYS	С	-15.51	-30.80	77.93	15.00
<b>77</b>	LYS	0	-15.07	-31.05	79.04	15.00
78	asn	N	-15.87	-29.58	77.55	15.00
78	ASN	CA	-15.78	-28.42	78.42	15.00
78	ASN	CB	-16.11	-27.17	77.61	15.00
78	asn	CG	-15.75	-25.90	78.33	15.00
78	ASN	OD1	-14.65	-25.77	78.86	15.00
78	ASN	ND2	-16.66	-24.93	78.33	15.00
78		C	-16.76	-28.55	79.61	15.00
78	ASN	0	-16.66	-27.82	80.59	15.00
79	ARG		-17.70	-29.48	79.47	15.00
79	ARG		-18.73	-29.76	80.46	15.00
79	ARG	CB	-18.11	-30.22	81.77	15.00
79	ARG	CG	-17.42	-31.56	81.70	15.00
79	ARG		-16.95	-31.93	83.08	15.00
79	ARG			-33.15	83.11	15.00
79	ARG		-14.86	-33.22	82.89	15.00
79	ARG	NH1	14.17	-32.13	82.58	15.00
79	ARG	NH2	-14.21	-34.37	33.07	15.00

79	ARG	. C	-19.66	-28.58	80.71	15.00
79	ARG	0	-20.27	-28.49	81.78	15.00
80	GLY	N	-19.79	-27.70	79.72	15.00
80	GLY	CA	-20.65	-26.55	79.87	15.00
80	GLY	C	-20.48	-25.48	78.81	15.00
80	GLY	0	-19.54	-25.52	78.01	15.00
81	ILE	N	-21.41	-24.53	78.80	15.00
81	ILE	CA	-21.40	-23.40	77.88	15.00
81	ILE	CB	-22.23	-23.68	76.58	15.00
81	ILE	CG2	-23.68	-24.05	76.93	15.00
81	ILE	CG1	-22.19	-22.46	75.64	15.00
81	ILE	CD1	-22.89	-22.66	74.31	15.00
81	ILE	C	-21.99	-22.20	78.62	15.00
81	ILE	0	-22.91	-22.35	79.43	15.00
82	ASP	N	-21.44	-21.02	78.37	15.00
82	ASP	CA	-21.91	-19.81	79.01	15.00
82	ASP	CB	-20.85	-18.71	78.94	15.00
82	ASP	CG	-19.73	-18.92	79.95	15.00
82	ASP	OD1	-18.66	-18.32	79.78	15.00
82	ASP	OD2	-19.93	-19.69	80.91	15.00
82	ASP	С	-23.21	-19.25	78.45	15.00
82	ASP	0	-23.63	-19.58	77.33	15.00
83	SER	N	-23.86	-18.43	79.26	15.00
83	SER	CA	-25.09	-17.78	78.85	15.00
83	SER	CB	-25.92	-17.36	80.08	15.00
83	SER	OG	-25.25	-16.38	80.86	15.00
83	SER	C	-24.65	-16.55	78.06	15.00
83	SER		-23.50	-16.10	78.17	15.00
84	GLU	N	-25.56	-16.01	77.26	15.00
84	GLU	CA.	-25.27	-14.83	76.47	15.00
84	GLU	CB	-26.53	-14.40	75.70	15.00
	GLU		-26.39	-13.10	74.93	15.00
84	GLU	CD	-25.30	-13.12	73.85	15.00
84	GLU	OE1	-24.76	-12.04	73.52	15.00
	GLU		-24.98	-14.22	73.33	15.00
	GLU	С	-24.77	-13.70	77.38	15.00
84			-23.77	-13.09	77.06	15.00
	ASP		-25.44	-13.47	78.51	15.00
	ASP		-25.05	-12.41	79.46	15.00
	ASP		-26.03	-12.35	80.65	15.00
	ASP		-27.20	-11.39	80.42	15.00
		OD1	-27.86		79.37	15.00
85	ASP	OD2	-27.47	-10.53	81.29	15.00

85	ASP	С	-23.62	-12.54	79.98	15.00
85	ASP	0	-22.89	-11.55	80.07	15.00
86	ALA	N	-23.23	-13.77	80.27	15.00
86	ALA	CA	-21.91	-14.09	80.78	15.00
86	ALA	CB	-21.95	-15.45	81.49	15.00
86	ALA	С	-20.79	-14.09	79.71	15.00
86	ALA	0	-19.61	-14.21	80.04	15.00
87	TYR	N	-21.16	-13.98	78.44	15.00
87	TYR	CA	-20.18	-13.99	77.36	15.00
87	TYR	CB	-19.75	-15.44	77.09	15.00
87	TYR	CG	-18.36	-15.67	76.52	15.00
87	TYR	CD1	-17.82	-14.83	75.54	15.00
87	TYR	CE1	-16.55	-15.09	74.99	15.00
87	TYR	CD2	-17.60	-16.76	76.94	15.00
87	TYR	CE2	-16.34	-17.03	76.40	15.00
87	TYR	CZ	-15.82	-16.19	75.42	15.00
87	TYR	OH	-14.59	-16.49	74.88	15.00
87	TYR	C	-20.90	-13.42	76.14	15.00
87	TYR	0	-21.25	-14.15	75.22	15.00
88	PRO	N	-21.09	-12.09	76.11	15.00
88	PRO	CD	-20.58	-11.10	77.08	15.00
88	PRO	CA	-21.77	-11.42	75.00	15.00
88	PRO	CB	-21.83	-9.97	75.47	15.00
88	PRO	CG	-20.62	-9.82	76.28	15.00
88	PRO	C	-21.13	-11.58	73.62	15.00
88	PRO	0	-19.92	-11.77	73.47	15.00
89	TYR	N	-21.98	-11.45	72.61	15.00
89	TYR	CA	-21.56	-11.59	71.22	15.00
89	TYR	CB	-22.77	-11.93	70.35	15.00
89	TYR	CG	-22.41	-12.26	68.92	15.00
89	TYR	CD1	-21.67	-13.41	68.62	15.00
89	TYR	CE1	-21.30	-13.71	67.32	15.00
89	TYR	CD2	-22.78	-11.43	67.87	15.00
89	TYR	CE2	-22.41	-11.72	66.56	15.00
89	TYR	CZ	-21.68	-12.87	66.30	15.00
89	TYR	OH	-21.29	-13.15	65.01	15.00
89	TYR	C	-20.91	-10.31	70.72	15.00
89	TYR	0	-21.45	-9.21	70.89	15.00
90	VAL	N	-19.73	-10.45	70.13	15.00
90	VAL	CA	-19.01	-9.31	69.58	15.00
90	VAL	CB	-17.59	-9.14	70.23	15.00
90	VAL	CG1	-17.71	-8.96	71.73	15.00
90	VAL	CG2	-16.69	-10.31	69.89	15.00

			<b>4</b> V	
90 VAL C	-18.88		68.06	15.00
90 VAL O	-18.50			15.00
91 GLY N	-19.17	-10.63		15.00
91 GLY CA	-19.08	-10.82	66.09	15.00
91 GLY C	-17.67	-10.74	65.55	15.00
91 GLY 0	-17.44	-10.33	64.40	15.00
92 GLN N	-16.71	-11.14	66.37	15.00
92 GLN CA	-15.31	-11.12	66.00	15.00
92 GLN CB	-14.73	-9.72	66.26	15.00
92 GLN CG	-13.35	-9.49	65.65	15.00
92 GLN CD	-12.71	-8.19	66.08	15.00
92 GLN OE1	-12.20	-7.44	65.24	15.00
92 GLN NE2	-12.73	-7.92	67.39	15.00
92 GLN C	-14.60	-12.19	66.82	15.00
92 GLN O	-15.02	-12.50	67.94	15.00
93 GLU N	-13.54	-12.78	66.26	15.00
93 GLU CA	-12.77	-13.82	66.94	15.00
93 GLU CB	-11.86	-14.56	65.94	15.00
93 GLU CG	-12.59	-15.03	64.68	15.00
93 GLU CD	-11.65	-15.50	63.57	15.00
93 GLU OE1	-10.57	-14.88	63.36	15.00
93 GLU OE2	-12.02	-16.49	62.88	15.00
93 GLU C	-11.92	-13.20	68.04	15.00
93 GLU O	-11.31	-12.14	67.84	15.00
94 GLU N	-11.90	-13.84	69.20	15.00
94 GLU CA	-11.15	-13.38	70.36	15.00
94 GLU CB	-12.03	-12.49	71.25	15.00
94 GLU CG	-12.71	-11.36	70.48	15.00
94 GLU CD	-13.30	-10.28	71.35	15.00
94 GLU OE1	-13.81	-10.60	72.43	15.00
94 GLU OE2	-13.27	-9.10	70.94	15.00
94 GLU C	-10.70	-14.63	71.11	15.00
94 GLU O	-11.13	-15.74	70.77	15.00
95 SER N	-9.82	-14.46	72.10	15.00
95 SER CA	-9.32	-15.60	72.86	15.00
95 SER CB	-8.13	-15.19	73.73	15.00
95 SER OG	-8.46	-14.11	74.58	15.00
95 SER C		-16.21	73.74	15.00
95 SER O		-15.51	74.26	15.00
96 CYS N		-17.52	73.95	15.00
96 CYS CA		-18.20	74.78	15.00
96 CYS C		-17.68	76.21	15.00
96 CYS O	-10.25	-17.71	76.89	15.00

96 CYS CB	-11.10	-19.72	74.74	15.00
96 CYS SG	-12.37	-20.66	75.65	
97 <b>MET</b> N	-12.43	-17.18	76.66	15.00
97 MET CA	-12.57	-16.63	77.99	15.00
97 MET CB	-12.71	-15.10	77.90	15.00
97 MET CG	-12.32	-14.33	79.15	15.00
97 MET SD	-10.53	-14.18	79.36	15.00
97 MET CE	-10.23	-15.44	80.61	15.00
97 MET C	-13.83	-17.23	78.62	15.00
97 MET O	-14.71	-16.50	79.08	15.00
98 TYR N	-13.95	-18.55	78.57	15.00
98 TYR CA	-15.11	-19.23	79.14	15.00
98 TYR CB	-15.04	-20.74	78.86	15.00
98 TYR CG	-16.04	-21.57	79.65	15.00
98 TYR CD1	-17.41	-21.44	79.44	15.00
98 TYR CE1	-18.32	-22.23	80.14	15.00
98 TYR CD2	-15.60	-22.52	80.59	15.00
98 TYR CE2	-16.51	-23.31	81.29	15.00
98 TYR CZ	-17.87	-23.16	81.06	15.00
98 TYR OH	-18.78	-23.95	81.71	15.00
98 TYR C	-15.16	-19.00	80.66	15.00
98 TYR O	-14.22	-19.33	81.36	15.00
99 ASN N	-16.23	-18.37	81.13	15.00
99 ASN CA	-16.39		82.56	15.00
99 ASN CB	-17.13	-16.81	82.81	15.00
99 ASN CG	-17.36	-16.54	84.30	15.00
99 ASN OD1	-17.39		85.12	15.00
99 ASN ND2	17.54	-15.27	84.65	15.00
99 ASN C	-17.18	-19.27	83.16	15.00
99 ASN O	-18.38	-19.39	82.93	15.00
100 PRO N	-16.53	-20.09	83.98	15.00
100 PRO CD	-15.13		84.42	15.00
100 PRO CA	-17.17	-21.24	84.63	15.00
100 PRO CB	-16.08	-21.75	85.57	15.00
100 PRO CG	-14.80		84.89	15.00
100 PRO C		-20.86	85.40	15.00
100 PRO O	-19.41	-21.59	85.40	15.00
101 THR N	-18.41	-19.69	86.04	15.00
101 THR CA	-19.54	-19.24	86.84	15.00
101 THR CB		-18.26	87.95	15.00
101 THR OG1		-17.04		15.00
101 THR CG2		-18.90	88.79	15.00
101 THR C	-20.73	-18.68	86.08	15.00

102 GLY N -20.57 -18.48 84.77 15.00 102 GLY CA -21.68 -17.98 83.97 15.00 102 GLY C -22.30 -19.14 83.21 15.00 102 GLY O -23.13 -18.94 82.32 15.00 103 LYS N -21.88 -20.35 83.54 15.00 103 LYS CA -22.36 -21.57 82.90 15.00 103 LYS CB -21.73 -22.79 83.58 15.00						
102 GLY CA		. 0	-21.79	-18.43	86.66	15.00
102 GLY CA		N	-20.57	-18.48		
102 GLY 0		CA	-21.68	-17.98	83.97	15.00
103 LYS N			-22.30	-19.14	83.21	15.00
103 LYS CA		0	-23.13	-18.94	82.32	15.00
103 LYS CB			-21.88	-20.35	83.54	15.00
103 LYS CG	103 LYS	CA	-22.36	-21.57	82.90	15.00
103 LYS CD			-21.73	-22.79	83.58	15.00
103 LYS CE				-24.13	82.98	15.00
103 LYS NZ				-25.25	83.81	15.00
103 LYS C		CE	-21.82	-25.07	85.28	15.00
103 LYS O			-21.22	-26.16	86.10	15.00
104 ALA N					82.95	15.00
104 ALA CA			-24.50	-21.54	84.00	15.00
104 ALA CB				-21.89	81.79	15.00
104 ALA C					81.72	15.00
104 ALA O					80.77	15.00
105 ALA N					81.32	15.00
105 ALA CA					81.46	15.00
105 ALA CB					80.78	15.00
105 ALA C					80.36	15.00
105 ALA O						15.00
106 LYS N						15.00
106 LYS CA						15.00
106 LYS CB						15.00
106 LYS CG						15.00
106 LYS CD					79.90	15.00
106 LYS CE						15.00
106 LYS NZ						
106 LYS C						
106 LYS O						
107 CYS N -24.03 -30.47 77.57 15.00 107 CYS CA -24.67 -31.55 76.85 15.00 107 CYS CB -24.96 -31.18 75.39 15.00 107 CYS SG -23.57 -31.18 74.26 15.00 107 CYS C -23.77 -32.77 76.95 15.00 107 CYS O -22.60 -32.67 77.31 15.00 108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
107 CYS CA -24.67 -31.55 76.85 15.00 107 CYS CB -24.96 -31.18 75.39 15.00 107 CYS SG -23.57 -31.18 74.26 15.00 107 CYS C -23.77 -32.77 76.95 15.00 107 CYS O -22.60 -32.67 77.31 15.00 108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
107 CYS CB -24.96 -31.18 75.39 15.00 107 CYS SG -23.57 -31.18 74.26 15.00 107 CYS C -23.77 -32.77 76.95 15.00 107 CYS O -22.60 -32.67 77.31 15.00 108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
107 CYS SG -23.57 -31.18 74.26 15.00 107 CYS C -23.77 -32.77 76.95 15.00 107 CYS O -22.60 -32.67 77.31 15.00 108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
107 CYS C -23.77 -32.77 76.95 15.00 107 CYS O -22.60 -32.67 77.31 15.00 108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
107 CYS O -22.60 -32.67 77.31 15.00 108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
108 ARG N -24.35 -33.94 76.70 15.00 108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
108 ARG CA -23.62 -35.19 76.78 15.00 108 ARG CB -24.30 -36.10 77.80 15.00						
108 ARG CB -24.30 -36.10 77.80 15.00						
108 ARG CG -24.56 -35.39 79.13 15.00						
	TOR ARG	CG	-24.56	-35.39	79.13	15.00

108 ARG CD	-24.78	-36.37	80.25	15.00
108 ARG NE	-26.08	-37.03		15.00
108 ARG CZ	-26.31	-38.29	80.55	15.00
108 ARG NH1	-25.34	-39.05	81.03	15.00
108 ARG NH2	-27.55	-38.77	80.48	15.00
108 ARG C	-23.52	-35.88	75.42	15.00
108 ARG O	-23.75		75.29	15.00
109 GLY N	-23.15	-35.11	74.40	15.00
109 GLY CA	-23.01	-35.66	73.07	15.00
109 GLY C	-23.96	-35.04	72.06	15.00
109 GLY 0	-24.53	-33.97	72.30	15.00
110 TYR N	-24.12	-35.71	70.92	15.00
110 TYR CA	-24.99	-35.26	69.85	15.00
110 TYR CB	-24.28	-34.21	68.99	15.00
110 TYR CG	-23.06	-34.73	68.24	15.00
110 TYR CD1	-21.77	-34.40	68.67	15.00
110 TYR CE1	-20.65	-34.80	67.94	15.00
110 TYR CD2	-23.19	-35.48	67.08	15.00
110 TYR CE2	-22.08	-35.89	66.35	15.00
110 TYR CZ	-20.81	-35.54	66.79	15.00
110 TYR OH	-19.71	-35.91	66.03	15.00
110 TYR C	-25.41	-36.44	68.98	15.00
110 TYR O	-24.81	-37.52	69.04	15.00
111 ARG N	-26.39	-36.20	68.12	15.00
111 ARG CA	-26.91	-37.23	67.23	15.00
111 ARG CB	-28.22	-37.78	67.83	15.00
111 ARG CG	-28.86	-38.97	67.11	15.00
ll1 ARG CD	-29.48	-38.54	65.78	15.00
lli arg ne	-30.34	-39.56	65.17	15.00
lll ARG CZ	-31.14	-39.34	64.13	15.00
lll arg NH1	-31.19	-38.13	63.58	15.00
111 ARG NH2	-31.92	-40.30	63.68	15.00
L11 ARG C	-27.13	-36.60	65.85	15.00
111 ARG O		-35.51	65.74	15.00
12 GLU N		-37.29		15.00
112 GLU CA	-26.82		63.43	15.00
112 GLU CB	-25.52	-37.09	62.65	15.00
112 GLU CG	-24.30	-36.39	63.23	15.00
12 GLU CD	-23.02	-36.62		15.00
.12 GLU OE1	-22.60		61. <b>69</b>	
.12 GLU OE2			62.55	
.12 GLU C	-27.99		62.76	15.00
.12 GLU O	-28.31	-38.67	63.08	15.00

113	ILE	N	-28.66	-36.80	61.85	15.00
113	ILE	CA	-29.79	-37.35	61.11	15.00
113	ILE	CB	-30.77	-36.21	60.69	15.00
113	ILE	CG2	-31.67	-36.67	59.56	15.00
113	ILE	CG1	-31.62	-35.77	61.89	15.00
113	ILE	CD1	-30.84	-35.23	63.06	15.00
113	ILE	С	-29.26	-38.07	59.87	15.00
113	ILE	0	-28.26	-37.65	59.27	15.00
114	PRO	N	-29.88	-39.21	59.49	15.00
114	PRO	CD	-31.02	-39.93	60.09	15.00
114	PRO	CA	-29.39	~39.90	58.30	15.00
114	PRO	СВ	-30.43	-41.00	58.08	15.00
114	PRO	CG	-30.91	-41.30	59.46	15.00
114	PRO	С	-29.36	-38.89	57.15	15.00
114	PRO	0	-30.36	-38.22	56.87	15.00
115	GLU :	N	-28.18	-38.70	56.57	15.00
115	GLU (	CA	-27.97	-37.75	55.49	15.00
115	GLU (	CB	-26.55	-37.89	54.96	15.00
115	GLU (	CG	-26.25	-37.14	53.69	15.00
115	GLU (	CD	-24.98	-37.66	53.04	15.00
115	GLU (	OE1	-25.08	-38.38	52.01	15.00
115	GLU (	OE2	-23.89	-37.37	53.58	15.00
115	GLU (	С	-28.99	-37.89	54.35	15.00
115	GLU (	0	-29.14	-38.97	53.79	15.00
116	GLY I	N	-29.66	-36.78	54.03	15.00
116	GLY (	CA	-30.66	-36.76	52.98	15.00
	GLY (	C	-32.05	-37.27	53.37	15.00
116	GLY (	0	-33.00	-37.16	52.59	15.00
117	ASN I	N	-32.17	-37.77	54.60	15.00
117	ASN (	CA	-33.43	-38.32	55.09	15.00
117	ASN (	CB	-33.14	-39.42	56.11	15.00
117	ASN (		-34.25	-40.47	56.18	15.00
117				-40.19	55.83	15.00
117	ASN 1	ND2	-33.91	-41.67	56.64	15.00
117	ASN (	C	-34.36	-37.28	55.72	15.00
	ASN (		-34.34	-37.10	56.93	15.00
	GLU 1			-36.64	54.89	15.00
118	GLU (	CA	-36.16	-35.64	55.36	15.00
	GLU (		-36.86	-34.94	54.19	15.00
	GLU (				53.50	15.00
	GLU (			-32.98		15.00
	GITA (				51.39	15.00
118	GLU (	DE2	-37.37	-31.97	53.11	15.00

118 GLU C	-37.21	-36.24	56.30	15.00
118 GLU O	-37.72	-35.57	57.19	15.00
119 LYS N	-37.53	-37.52	56.08	15.00
119 LYS CA	-38.47	-38.23	56.93	15.00
119 LYS CB	-38.64	-39.66	56.41	15.00
119 LYS CG	-39.73	-39.83	55.38	15.00
119 LYS CD	-39.53	-41.10	54.58	15.00
119 LYS CE	-38.51	-40.88	53.46	15.00
119 LYS NZ	-38.97	-39.83	52.48	15.00
119 LYS C	-37.91	-38.29	58.36	15.00
119 LYS O	-38.58		59.32	15.00
120 ALA N	-36.70		58.51	15.00
120 ALA CA	-36.06		59.81	15.00
120 ALA CB	-34.71	-39.60	59.68	15.00
120 ALA C	-35.91	-37.54	60.41	15.00
120 ALA O	-36.06	-37.38	61.61	15.00
121 LEU N	-35.63		59.56	15.00
121 LEU CA	-35.48		60.05	15.00
121 LEU CB	-35.11	-34.21	58.90	15.00
121 LEU CG	-34.87	-32.75	59.33	15.00
121 LEU CD1	-33.74	-32.69	60.38	15.00
121 LEU CD2	-34.53	-31.88	58.12	15.00
121 LEU C	-36.78		60.72	15.00
121 LEU O	-36.75	-34.17	61.83	15.00
122 LYS N	-37.91	-34.94	60.05	15.00
122 LYS CA	-39.23	-34.59	60.56	15.00
122 LYS CB	-40.31	-35.02	59.58	15.00
122 LYS CG	-41.74	-34.91	60.12	15.00
122 LYS CD	-42.72	-35.57	59.17	15.00
122 LYS CE	-44.11	-35.67	59.76	15.00
122 LYS NZ	-45.10	-36.14	58.74	15.00
122 LYS C	-39.44	-35.32	61.88	15.00
122 LYS O	-39.88	-34.74	62.87	15.00
123 ARG N	-39.14	-36.61	61.86	15.00
123 ARG CA	-39.28	-37.46	63.03	15.00
123 ARG CB	~38.80	-38.87	62.70	15.00
123 ARG CG	-38.84	-39.87	63.86	15.00
123 ARG CD	-37.70	-40.88	63.77	15.00
123 ARG NE	-37.37	-41.20	62.38	15.00
123 ARG CZ	-36.78	-42.32	61.98	
123 ARG NH1	-36.45		62.87	
123 ARG NH2	-36.53	-42.51		
123 ARG C		-36.88		

			-	
123 ARG O	-38.93	-36.84	65.32	15.00
124 ALA N	-37.25	-36.42	63.89	15.00
124 ALA CA	-36.37	-35.84	64.90	15.00
124 ALA CB	-34.98	-35.61	64.35	15.00
124 ALA C	-36.95	-34.53	65.45	15.00
124 ALA O	-37.02	-34.33	66.66	15.00
125 VAL N	-37.41	-33.66	64.57	15.00
125 VAL CA	-37.99	-32.41	65.03	15.00
125 VAL CB	-38.35	-31.48	63.87	15.00
125 VAL CG1	-38.98	-30.19	64.39	15.00
125 VAL CG2	-37.09	-31.17	63.06	15.00
125 VAL C	-39.23	-32.65	65.89	15.00
125 VAL 0	-39.46	-31.94	66.87	15.00
126 ALA N	-40.01	-33.67	65.53	15.00
126 ALA CA	-41.22	-34.03	66.25	15.00
126 ALA CB	-42.03	-35.02	65.44	15.00
126 ALA C	-40.89	-34.61	67.62	15.00
126 ALA 0	-41.40	-34.14	68.64	15.00
127 ARG N	-40.03	-35.62	67.64	15.00
127 ARG CA	-39.61	-36.29	68.86	15.00
127 ARG CB	-38.83	-37.54	68.48	15.00
127 ARG CG	-38.45	-38.44	69.64	15.00
127 ARG CD	-39.63	~39.28	70.13	15.00
127 ARG NE	-40.15	-40.18	69.10	15.00
127 ARG CZ	-41.18	-39.90	68.31	15.00
127 ARG NH1	-41.59	-40.78	67.41	15.00
127 ARG NH2	-41.81	-38.73	68.42	15.00
127 ARG C	-38.75	-35.47	69.83	15.00
127 ARG O	~39.05	-35.36	71.01	15.00
128 VAL N	-37.64	-34.94	69.31	15.00
128 VAL CA	-36.66	-34.21	70.11	15.00
128 VAL CB	-35.23	-34.54	69.59	15.00
128 VAL CG1	-34.17	-33.72	70.32	15.00
128 VAL CG2	-34.95	-36.02	69.74	15.00
128 VAL C	-36.82	-32.70	70.26	15.00
128 VAL O	-36.73	-32.17	71.36	15.00
129 GLY N	-37.00	-32.01	69.14	15.00
129 GLY CA	-37.16	-30.56	69.20	15.00
129 GLY C	-36.36	-29.91	68.08	15.00
129 GLY O	-35.89	-30.62	67.18	15.00
130 PRO N	-36.19	-28.57	68.11	15.00
130 PRO CD	-36.66	-27.55	69.17	15.00
130 PRO CA	-35.45	-27.83	57.09	15.00

130	PRO	CB	-35.23	-26.47	67.74	15.00
130	PRO	CG	-36.51	-26.30	68.52	15.00
130	PRO	С	-34.13	-28.52	66.74	15.00
130	PRO	0	-33.37	-28.91	67.63	15.00
131	VAL	N	-33.90	-28.72	65.45	15.00
131	VAL	CA	-32.71	-29.40	64.97	15.00
131	VAL	CB	-33.13	-30.65	64.15	15.00
131	VAL	CG1	-31.93	-31.36	63.57	15.00
131	VAL	CG2	-33.92	-31.61	65.03	15.00
131	VAL	С	-31.82	-28.49	64.13	15.00
131	VAL	0	-32.32	-27.69	63.34	15.00
132	SER	N	-30.51	-28.59	64.32	15.00
132	SER	CA	-29.56	-27.79	63.55	15.00
132	SER	CB	-28.18	-27.83	64.22	15.00
132	SER	OG	-28.20	-27.34	65.55	15.00
132	SER	C	-29.44	-28.37	62.14	15.00
132	SER	0	-29.41	-29.59	61.96	15.00
133	VAL	N	-29.39	-27.50	61.13	15.00
133	VAL	CA	-29.25	-27.94	59.75	15.00
133	VAL	CB	-30.63	-28.07	59.03	15.00
133	VAL	CG1	-31.40	-29.29	59.53	15.00
133	VAL	CG2	-31.45	-26.80	59.20	15.00
133	VAL	С	-28.37	-27.00	58.94	15.00
133	VAL	0	-28.12	-25.87	59.34	15.00
134	ALA	N	-27.86	-27.50	57.82	15.00
134	ALA	CA	-27.03	-26.73	56.91	15.00
	ALA		-25.65	-27.34	56.76	15.00
	ALA		-27.75	-26.73	55.57	15.00
	ALA		-28.29	-27.76	55.14	15.00
	ILE	N	-27.80	-25.58	54.94	15.00
135	ILE		-28.49	-25.45	53.66	15.00
	ILE		-29.82	-24.68	53.82	15.00
	ILE		-30.80	-25.45	54.70	15.00
	ILE		-29.52	-23.28	54.38	15.00
		CD1	-30.72		54.52	15.00
135	ILE	С	-27.64	-24.61	52.73	15.00
135		0	-26.59	-24.09	53.13	15.00
	ASP		-28.09	-24.49	51.49	15.00
	ASP			-23.66	50.51	15.00
	ASP			-24.28	49.11	15.00
	ASP			-23.32	48.02	15.00
	ASP			-23.49	46.88	15.00
136	ASP	OD2	-26.22	-22.40	48.29	15.00

136 ASP C	-28.19	-22.36	50.54	15.00
136 ASP 0	-29.34		50.10	15.00
137 ALA N	-27.58		51.11	15.00
137 ALA CA	-28.19		51.21	15.00
137 ALA CB	-28.14	-19.54	52.65	15.00
137 ALA C	-27.47		50.30	15.00
137 ALA O	-27.42	-17.85	50.60	15.00
138 SER N	-26.94	-19.53	49.20	15.00
138 SER CA	-26.20		48.30	15.00
138 SER CB	-25.12	-19.44	47.57	15.00
138 SER OG	-25.71		46.78	15.00
138 SER C	-27.06	-17.91	47.29	15.00
138 SER O	-26.76	-16.77	46.94	15.00
139 LEU N	-28.12	-18.55	46.82	15.00
139 LEU CA	-28.99		45.82	15.00
139 LEU CB	-30.12		45.46	15.00
139 LEU CG	-29.76	-20.02	44.48	15.00
139 LEU CD1	-29.30	-19.40	43.18	15.00
139 LEU CD2	-28.67	-20.89	45.03	15.00
139 LEU C	-29.56	-16.58	46.18	15.00
139 LEU O	-29.93	-16.32	47.33	15.00
140 THR N	-29.63	-15.71	45.19	15.00
140 THR CA	-30.19	-14.37	45.38	15.00
140 THR CB	-30.06	-13.54	44.09	15.00
140 THR OG1	-28.67	-13.30	43.82	15.00
140 THR CG2	-30.80	-12.20	44.20	15.00
140 THR C	-31.65	-14.43	45.86	15.00
140 THR O	-32.09	-13.62	46.69	15.00
141 SER N	-32.38	-15.45	45.40	15.00
141 SER CA	-33.77	-15.63	45.79	15.00
141 SER CB	-34.39	-16.77	44.99	15.00
141 SER OG	-33.78	-18.00	45.32	15.00
141 SER C	-33.89	-15.89	47.29	15.00
141 SER O	-34.94	-15.63	47.90	15.00
142 PHE N		-16.43	47.89	15.00
L42 PHE CA	-32.79	-16.70	49.33	15.00
142 PHE CB	-31.71	-17.72	49.67	15.00
142 PHE CG	-31.77	-18.20	51.08	15.00
142 PHE CD1	-32.45	-19.38	51.39	15.00
42 PHE CD2	-31.14	-17.49	52.10	15.00
42 PHE CE1	-32.51	-19.85	52.70	15.00
.42 PHE CE2	-31.20	-17.95	53.41	15.00
.42 PHE CZ	-31.88	-19.14	53.72	15.00

142	PHE	С	-32.53	-15.40	50.08	15.00
142	PHE	0	-33.22	-15.09	51.06	15.00
143	GLN	N	-31.55	-14.64	49.60	15.00
143	GLN	CA	-31.19	-13.38	50.22	15.00
143	GLN	CB	-30.07	-12.72	49.44	15.00
143	GLN	CG	-29.68	-11.32	49.94	15.00
143	GLN	CD	-28.36	-10.81	49.37	15.00
143	GLN	OE1	-28.05	-9.62	49.47	15.00
143	GLN	NE2	-27.56	-11.71	48.80	15.00
143	GLN	С	-32.38	-12.43	50.33	15.00
143	GLN	0	-32.57	-11.75	51.35	15.00
144	PHE	N	-33.22	-12.40	49.30	15.00
144	PHE	CA	-34.36	-11.50	49.31	15.00
144	PHE	CB	-34.41	~10.73	47.98	15.00
144	PHE	CG	-33.22	-9.85	47.75	15.00
144	PHE	CD1	-33.13	-8.59	48.35	15.00
144	PHE	CD2	-32.17	-10.27	46.94	15.00
144	PHE	CE1	-32.02	-7.77	48.15	15.00
144	PHE	CE2	-31.05	-9.46	46.73	15.00
144	PHE		-30.98	-8.21	47.34	15.00
144	PHE	С	-35.73	-12.13	49.64	15.00
144	PHE		-36.77	-11.52	49.38	15.00
	TYR		-35.72	-13.30	50.26	15.00
145			-36.97	-13.98	50.63	15.00
145	TYR		-36.68	-15.35	51.26	15.00
145	TYR		-37.89	-15.98	51.94	15.00
145	TYR		-38.72	-16.89	51.26	15.00
145	TYR		-39.83	-17.45	51.89	15.00
145	TYR		-38.21	-15.66	53.26	15.00
145	TYR		-39.33	-16.22	53.90	15.00
145	TYR		-40.13	-17.12	53.21	15.00
145			-41.21	-17.68	53.85	15.00
145	TYR		-37.81	-13.16	51.62	15.00
145	TYR		-37.27	-12.49	52.50	15.00
146			-39.13	-13.26	51.50	15.00
146	SER		-40.03	-12.55	52.41	15.00
146	SER		-40.33	-11.12	51.96	15.00
146	SER		-40.50	-11.03	50.57	15.00
146	SER		-41.31	-13.30	52.73	15.00
146	SER		-41.74	-13.31	53.88	15.00
147	LYS	N	-41.89	-14.00	51.76	15.00
147		CA	-43.13	-14.74	51.99	15.00
147	LYS	CB	-44.34	-13.80	51.82	15.00

147	LYS	CG	-44.60	-12.90	53.05	15.00
147	LYS	CD	-45.56	-11.75	52.75	15.00
147	LYS	CE	-45.63	-10.76	53.91	15.00
147	LYS	NZ	-46.25	-11.32	55.15	15.00
147	LYS	С	-43.30	-15.99	51.13	15.00
147	LYS	0	-42.68	-16.13	50.07	15.00
148	GLY	N	-44.12	-16.92	51.60	15.00
148	GLY	CA	-44.36	-18.14	50.87	15.00
148	GLY	С	-43.35	-19.22	51.18	15.00
148	GLY	0	-42.39	-19.00	51.90	15.00
149	VAL	N	-43.60	-20.42	50.67	15.00
149	VAL	CA	-42.69	-21.53	50.90	15.00
149	VAL	CB	-43.43	-22.89	50.72	15.00
149	VAL	CG1	-42.46	-24.05	50.79	15.00
149	VAL	CG2	-44.49	-23.05	51.80	15.00
149	VAL	С	-41.56	-21.34	49.88	15.00
	VAL	0	-41.81	-21.30	48.67	15.00
	TYR	N	-40.34	-21.13	50.37	15.00
150	TYR	CA	-39.19	-20.92	49.50	15.00
150	TYR	CB	-37.99	-20.42	50.30	15.00
150			-36.72	-20.27	49.46	15.00
	TYR		-36.48	-19.11	48.72	15.00
	TYR		-35.31	-18.98	47.96	15.00
150	TYR		-35.77	-21.30	49.41	15.00
150	TYR		-34.59	-21.17	48.65	15.00
150	TYR		-34.37	-20.01	47.93	15.00
150	TYR		-33.22	-19.88	47.19	15.00
150	TYR		-38.80	-22.18	48.76	15.00
150	TYR		-38.87	-23.29	49.30	15.00
151	TYR			-21.98	47.54	15.00
151	TYR		-37.87	-23.07	46.68	15.00
	TYR		-39.07	-23.95	46.31	15.00
151	TYR		-38.80	-25.01	45.27	15.00
	TYR		-38.36		45.62	15.00
	TYR		-38.19		44.66	15.00
	TYR			-24.75	43.92	15.00
	TYR		-38.88		42.96	15.00
151				-26.98	43.33	15.00
151	TYR			-27.95	42.36	15.00
	TYR			-22.44	45.44	15.00
	TYR			-21.37	45.00	15.00
	ASP			-23.11		
152	ASP	CA	-35.52	-22.62	43.70	15.00

152	ASP	CB	-34.55	-21.50	44.08	15.00
152	ASP	CG	-33.97	-20.77	42.87	15.00
152	ASP	OD1	-33.75	-21.39	41.81	15.00
152	ASP	OD2	-33.72	-19.56	42.99	15.00
152	ASP	С	-34.75	-23.77	43.07	15.00
152	ASP	0	-33.90	-24.38	43.73	15.00
153	GLU	N	-35.01	-24.02		15.00
153	GLU	CA	-34.35	-25.09	41.04	15.00
153	GLU	CB	-34.78	-25.06	39.56	15.00
153	GLU	CG	-36.22	-25.45	39.26	15.00
153	GLU	CD	-36.59	-25.21	37.78	15.00
153	GLU	OE1	-37.05		37.45	15.00
153	GLU	OE2	-36.40	-26.14	36.96	15.00
153	GLU	С	-32.84	-24.95	41.09	15.00
153	GLU	0	-32.12	-25.93	40.99	15.00
154	SER	N	-32.38	-23.71	41.20	15.00
154	SER	CA	-30.95	-23.41	41.24	15.00
154	SER	CB	-30.73	-21.93	40.92	15.00
154	SER	OG	-31.33	-21.57	39.69	15.00
154	SER	С	-30.25	-23.78	42.55	15.00
154	SER	o	-29.02	-23.75	42.63	15.00
155	CYS :	N	-31.02	-24.11	43.58	15.00
155	CYS	CA	-30.44	-24.46	44.87	15.00
155	CYS	С	-29.58	-25.72	44.76	15.00
155	CYS	0	-29.98	-26.72	44.17	15.00
155	CYS	CB	-31.53	-24.65	45.92	15.00
155	CYS	SG	-31.12	-23.81	47.48	15.00
156	ASN :	N	-28.40	-25.67	45.35	15.00
156	ASN	CA	-27.48	-26.80	45.29	15.00
156	asn (	CB	-26.09	-26.28	44.91	15.00
156	ASN (	CG	-25.15	-27.39	44.45	15.00
156	ASN (	OD1	-25.21	-28.54	44.94	15.00
	ASN I		-24.26	-27.05	43.53	15.00
156	ASN (	С	-27.42	-27.58	46.61	15.00
	ASN (		-26.99	-27.06	47.64	15.00
	SER I		-27.80	-28.85	46.54	15.00
157	SER (	CA	-27.82	-29.72	47.71	15.00
157	SER (	CB	-28.66	-30.98	47.45	15.00
157	SER (	OG	-28.07	-31.81	46.47	15.00
	SER (		-26.43	-30.14	48.17	15.00
	SER (		-26.27	-30.76	49.23	15.00
1.58	ASP 1	:1	-25.42	-29.82	47.37	15.00
158 .	ASP (	CA	-24.06	-30.18	47.69	15.00

158	ASP	CB	-23.44	-30.99	46.55	15.00
158	ASP	CG	-23.70	-32.49	46.69	15.00
158	ASP	OD1	-24.30	-33.11	45.78	15.00
158	ASP	OD2	-23.32	-33.05	47.75	15.00
158	ASP	С	-23.21	-29.01	48.14	15.00
158	ASP	0	-22.19	-29.21	48.79	15.00
159	ASN	N	-23.67	-27.80	47.85	15.00
159	ASN	CA	-22.95	-26.60	48.28	15.00
159	ASN	CB	-23.04	-25.49	47.22	15.00
159	ASN	CG	-22.27	-24.23	47.61	15.00
159	ASN	OD1	-21.81	-24.09	48.74	15.00
159	ASN	ND2	-22.14	-23.31	46.67	15.00
159	ASN	С	-23.55	-26.10	49.61	15.00
159	ASN	0	-24.20	-25.06	49.67	15.00
160	LEU	N	-23.38	-26.89	50.68	15.00
160	LEU	CA	-23.91	-26.47	51.98	15.00
160	LEU	CB	-23.83	-27.61	52.99	15.00
160	LEU	CG	-24.49	-28.94	52.62	15.00
160	LEU	CD1	-24.41	-29.89	53.81	15.00
160	LEU	CD2	-25.94	-28.71	52.23	15.00
160	LEU		-23.04	-25.30	52.41	15.00
160	LEU		-21.82	-25.42	52.46	15.00
161			-23.65	-24.15	52.70	15.00
161	ASN		-22.86	-22.98	53.07	15.00
161	ASN		-22.59	-22.13	51.83	15.00
161	ASN		-23.82	-21.92	50.98	15.00
161	ASN		-24.77	-21.24	51.38	15.00
161	ASN	ND2	-23.83	-22.51	49.81	15.00
161	ASN		-23.40	-22.10	54.20	15.00
161	ASN		-22.80	-21.07	54.54	15.00
162	HIS		-24.47	-22.55	54.85	15.00
162	HIS		-25.07	-21.73	55.92	15.00
162	HIS	CB	-26.12	-20.83	55.33	15.00
162	HIS		-26.58 -27.81	-19.75	56.27	15.00
162	HIS			-19.22	56.48	15.00
162	HIS		-25.73	-19.08	57.12	15.00
162			-26.41	-18.20	57.83	15.00
162	HIS		-27.68 -25.71	-18.27 -22.7€	57.46 56.91	15.00 15.00
162	HIS					15.00
162	HIS			-23.71	56.50	
163	ALA			-22.58	58.19	15.00
163	ALA			~23.43	59.24	15.00
163	ALA	CB	-24.95	-23.66	60.35	15.00

163	ALA	C	-27.17	-22.65	59.74	15.00
163	ALA	0	-27.10	-21.42	59.92	15.00
164	VAL	N	-28.27	-23.34	59.96	15.00
164	VAL	CA	-29.51	-22.71	60.36	15.00
164	VAL	CB	-30.30	-22.38	59.08	15.00
164	VAL	CG1	-31.28	-23.48	58.75	15.00
164	VAL	CG2	-30.90	-20.99	59.15	15.00
164	VAL	С	-30.30	-23.60	61.34	15.00
164	VAL	0	-29.89	-24.73	61.62	15.00
165	LEU	N	-31.40	-23.09	61.89	15.00
165	LEU	CA	-32.18	-23.86	62.86	15.00
165	LEU	CB	-32.27	-23.11	64.20	15.00
165	LEU	CG	-32.79	-23.87	65.44	15.00
165	LEU	CD1	-31.76	-24.89	65.90	15.00
165	LEU	CD2	-33.13	-22.90	66.57	15.00
165	LEU	С	-33.58	-24.27	62.40	15.00
165	LEU	0	-34.39	-23.42	62.03	15.00
166	ALA	N	-33.85	-25.57	62.43	15.00
166	ALA	CA	-35.15	-26.11	62.05	15.00
166	ALA	CB	-35.00	-27.51	61.50	15.00
166	ALA	C	-35.99	-26.11	63.33	15.00
166	ALA	0	-35.76	-26.91	64.25	15.00
167	VAL	N	-36.94	-25.19	63.39	15.00
167	VAL	CA	-37.80	-25.06	64.55	15.00
167	VAL	CB	-37.81	-23.60	65.03	15.00
167	VAL	CG1	-38.83	-22.75	64.24	15.00
167	VAL	CG2	-38.06	-23.55	66.50	15.00
167	VAL	C	-39.23	-25.58	64.33	15.00
167	VAL	0	-40.15	-25.31	65.14	15.00
168	GLY	N	-39.44	-26.31	63.24	15.00
168	GLY	CA	-40.76	-26.84	62.97	15.00
168	GLY		-40.97	-27.25	61.53	15.00
	GLY		-40.02	-27.37	60.74	15.00
169	TYR	N	-42.23	-27.48	61.20	15.00
	TYR			-27.89	59.87	15.00
169	TYR		-42.21	-29.33	59.54	15.00
169	TYR	CG	-42.75	-30.41	60.46	15.00
169	TYR		-43.94	-31.09	60.16	15.00
169	TYR	CE1.	-44.43	-32.10	60.99	15.00
169	TYR		-42.06	-30.79	61.61	15.00
169	TYR	CE2	-42.53		62.45	15.00
169	TYR	CZ		-32.45	62.13	15.00
169	TYR	OH	-44.15	-33.47	62.94	15.00

169	TYR	C	-44.18	-27.80	59.84	15.00
169	TYR	0	-44.84	-27.82	60.88	15.00
170	GLY	N	-44.73	-27.73	58.64	15.00
170	GLY	CA	-46.17	-27.63	58.48	15.00
170	GLY	С	-46.53	-27.64	57.01	15.00
170	GLY	0	-45.73	-28.04	56.18	15.00
171	ILE	N	-47.71	-27.14	56.69	15.00
171	ILE	CA	-48.16	-27.09	55.30	15.00
171	ILE	CB	-48.95	-28.37	54.91	15.00
171	ILE	CG2	-50.04	-28.69	55.91	15.00
171	ILE	CG1	-49.52	-28.25	53.51	15.00
171	ILE	CD1	-50.20	-29.53	53.05	15.00
171	ILE	C	-48.97	-25.82	55.09	15.00
171	ILE	0	-49.84	-25.48	55.89	15.00
172	GLN	N	-48.59	-25.07	54.05	15.00
172	GLN	CA	-49.21	-23.80	53.72	15.00
172	GLN	CB	-48.19	-22.67	53.89	15.00
172	GLN	CG	-48.68	-21.32	53.45	15.00
172	GLN	CD	-47.73	-20.20	53.84	15.00
172	GLN	OE1	-47.84	-19.61	54.92	15.00
172	GLN	NE2	-46.78	-19.89	52.96	15.00
172	GLN	С	-49.73	-23.81	52.29	15.00
172	GLN	0	-48.96	-23.96	51.33	15.00
	LYS	N	-51.04	-23.64	52.17	15.00
	LYS	CA	-51.73	-23.62	50.88	15.00
173	LYS	CB	-51.37	-22.35	50.09	15.00
173	LYS	CG	-51.48	-21.01	50.88	15.00
	LYS	CD	-52.85	-20.77	51.55	15.00
173	LYS	CE	-54.02	-20.84	50.56	15.00
	LYS	NZ	-53.89	-19.90	49.41	15.00
	LYS	С	-51.39	-24.87	50.08	15.00
		0	-51.27	-24.82	48.87	15.00
	GLY :		-51.22	-25.99	50.77	15.00
	GLY		-50.91	-27.24	50.10	15.00
	GLY		-49.43	-27.58	50.06	15.00
	GLY (		-49.07	-28.71	49.74	15.00
	ASN !		-48.59	-26.65	50.48	15.00
	ASN (		-47.15	-26.87	50.44	15.00
	asn (		-46.44	-25.64	49.88	15.00
	ASN (		-47.06	-25.14	48.59	15.00
	ASN (		-47.08	-25.86	47.58	15.00
	ASN 1		-47.56	-23.91	48.62	15.00
175 .	ASN (	С	-46.54	-27.23	51.79	15.00

175	ASN	0	-46.63	-26.46	52.74	15.00
176	LYS	N	-45.92	-28.40	51.87	15.00
176	LYS	CA	-45.28	-28.84	53.09	15.00
176	LYS	CB	-44.98	~30.33	53.02	15.00
176	LYS	CG	-46.23	-31.18	52.84	15.00
176	LYS	CD	-45.95	-32.63	53.09	15.00
176	LYS	CE	-47.17	-33.48	52.82	15.00
176	LYS	NZ	-46.84	-34.93	52.78	15.00
176	LYS	С	-44.00	-28.03	53.20	15.00
176	LYS	0	-43.37	-27.73	52.19	15.00
177	HIS	N	-43.59	-27.69	54.42	15.00
177	HIS	CA	-42.39	-26.88	54.58	15.00
177	HIS	CB	-42.76	-25.39	54.51	15.00
177	HIS	CG	-43.62	-24.94	55.64	15.00
177	HIS	CD2	-43.32	-24.68	56.94	15.00
177	HIS	ND1	-44.96	-24.70	55.50	15.00
177	HIS	CE1	-45.46	-24.31	56.66	15.00
177	HIS	NE2	-44.48	-24.29	57.55	15.00
177	HIS	С	-41.68	-27.10	55.88	15.00
177	HIS	0	-42.20	-27.76	56.78	15.00
178	TRP	N	-40.52	-26.46	55.99	15.00
178	TRP	CA	-39.69	-26.51	57.18	15.00
178	TRP	CB	-38.24	-26.84	56.79	15.00
178	TRP	CG	-38.01	-28.23	56.31	15.00
178	TRP	CD2	-38.14	-29.45	57.06	15.00
178	TRP	CE2	-37.73	-30.50	56.22	15.00
	TRP	CE3	-38.56	-29.74	58.37	15.00
178	TRP	CD1	-37.55	-28.59	55.07	15.00
178	TRP	NEl	-37.38	-29.95	55.01	15.00
178	TRP	CZ2	-37.73	-31.85	56.64	15.00
178	TRP	CZ3	-38.56	-31.08	58.79	15.00
	TRP		-38.14	-32.12	57.92	15.00
	TRP		-39.73	-25.11	57.74	15.00
178	TRP	0	-39.68	-24.14	56.97	15.00
179	ILE		-39.90	-24.97	59.05	15.00
179	ILE		-39.91	-23.65	59.65	15.00
179	ILE	CB	-40.89	-23.55	60.84	15.00
179	ILE		-40.95	-22.10	61.34	15.00
179	ILE	CG1	-42.29	-24.02	60.43	15.00
179	ILE		-43.32	-23.95	61.55	15.00
179		С	-38.47	-23.41	60.08	15.00
179	ILE		-37.96	-24.05	61.00	15.00
180	ILE	N	-37.79	-22.52	59.36	15.00

180 ILE CA	-36.40	-22.23	59.61	15.00
180 ILE CB		-22.34	58.29	15.00
180 ILE CG	2 -34.14	-21.97	58.52	15.00
180 ILE CG	1 -35.74	-23.74	57.68	15.00
180 ILE CD	1 -35.52	-24.88	58.66	15.00
180 ILE C	-36.13	-20.88	60.25	15.00
180 ILE O	-36.68	-19.85	59.84	15.00
181 LYS N	-35.24	-20.90	61.25	15.00
181 LYS CA	-34.84	-19.72	61.99	15.00
181 LYS CB	-34.77	-20.06	63.48	15.00
181 LYS CG	-34.66	-18.84	64.35	15.00
181 LYS CD	-34.34	-19.17	65.79	15.00
181 LYS CE	-34.16	-17.89	66.56	15.00
181 LYS NZ	-33.69	-18.09	67.94	15.00
181 LYS C	-33.46	-19.27	61.49	15.00
181 LYS O	-32.50	-20.03	61.56	15.00
182 ASN N	-33.37	-18.05	60.98	15.00
182 ASN CA	-32.11	-17.52	60.47	15.00
182 ASN CB	-32.33	-16.76	59.15	15.00
182 ASN CG	-31.05	-16.64	58.29	15.00
182 ASN OD1	-30.00	-17.21	58.60	15.00
182 ASN ND2	-31.15	-15.91	57.18	15.00
182 ASN C	-31.43	-16.62	61.51	15.00
182 ASN 0	-32.00	-16.33	62.57	15.00
183 SER N	-30.18	-16.25	61.24	15.00
183 SER CA	-29.43	-15.38	62.14	15.00
183 SER CB	-28.25	-16.13	62.75	15.00
183 SER OG	-27.48	-16.78	61.76	15.00
183 SER C	-28.96	-14.09	61.44	15.00
183 SER O	-27.85	-13.63	61.66	15.00
184 TRP N	-29.82	-13.52	60.59	15.00
184 TRP CA	-29.50	-12.28	59.88	15.00
184 TRP CB	-29.69	-12.45	58.37	15.00
184 TRP CG		-13.44	57.76	15.00
184 TRP CD2		-13.98	56.44	15.00
184 TRP CE2		-14.83	56.30	15.00
184 TRP CE3		-13.83	55.34	15.00
184 TRP CD1	-27.60	-13.98	58.37	15.00
184 TRP NE1	-26.95	-14.81	57.50	15.00
184 TRP CZ2		-15.53	55.13	15.00
184 TRP CZ3		-14.52	54.17	15.00
184 TRP CH2		-15.35	54.07	15.00
184 TRP C	-30.35	-11.11	50.40	15.00

184	TRP	0	-30.51	-10.09	59.72	15.00
185	GLY	N	-30.84	-11.25	61.63	15.00
185	GLY	CA	-31.67	-10.22	62.24	15.00
185	GLY	С	-33.14	-10.43	61.94	15.00
185	GLY	0	-33.49	-11.18	61.03	15.00
186	GLU	N	-33.99	-9.73	62.68	15.00
186	GLU	CA	-35.45	-9.83	62.50	15.00
186	GLU	CB	-36.17	-9.31	63.74	15.00
186	GLU	CG	-35.94	-10.14	64.97	15.00
186	GLU	CD	-37.15	-10.15	65.89	15.00
186	GLU	OE1	-38.26	-10.52	65.41	15.00
186	GLU	OE2	-37.01	-9.81	67.09	15.00
186	GLU	С	-36.05	-9.15	61.27	15.00
186	GLU	0	-37.10	-9.56	60.78	15.00
187	asn	N	-35.38	-8.12	60.76	15.00
187	ASN	CA	-35.89	-7.42	59.59	15.00
187	asn	CB	-35.32	-6.00	59.52	15.00
187	asn	CG	-35.95	-5.04	60.53	15.00
187	ASN	OD1	-35.82	-3.83	60.39	15.00
187	ASN	ND2	-36.62	-5.58	61.56	15.00
187	ASN	С	-35.60	-8.14	58.29	15.00
187	asn	0	-35.82	-7.57	57.22	15.00
188	TRP	N	-35.12	-9.38	58.37	15.00
188	TRP	CA	-34.81	-10.18	57.17	15.00
188	TRP	CB	-33.42	-10.84	57.28	15.00
188	TRP	CG	-33.16	-11.85	56.18	15.00
188	TRP		-33.54	-13.23	56.17	15.00
188	TRP	CE2	-33.19	-13.75	54.89	15.00
188	TRP		-34.14	-14.09	57.10	15.00
188	TRP		-32.60	-11.59	54.97	15.00
188	TRP	NE1	-32.63	-12.73	54.18	15.00
188	TRP		-33.44	-15.08	54.52	15.00
188	TRP		-34.39	-15.42	56.74	15.00
188	TRP		-34.03	-15.90	55.46	15.00
	TRP		-35.88	-11.27	57.03	15.00
188	TRP		-36.45	-11.71	58.04	15.00
		N	-36.14	-11.69	55.79	15.00
189	GLY		-37.13	-12.72	55.53	15.00
189	GLY		-38.45	-12.40	56.21	15.00
189	GLY		-38.91	-11.26	56.15	15.00
	ASN		-39.07	-13.38	56.85	15.00
190	asn		-40.33	-13.14	57.55	15.00
190	ASN	CB	- 41 33	-14.27	57.29	15.00

	) ASN		-42.74	-13.89	57.69	15.00
190		OD1	-42.98	-12.88	58.35	15.00
190	ASN	ND2	-43.69	-14.70	57.26	15.00
190	ASN	C	-40.03	-13.04	59.04	15.00
190	ASN	0	-40.09	-14.04	59.76	15.00
191	LYS	N	-39.68	-11.83	59.49	15.00
191	LYS	CA	-39.34	-11.59	60.90	15.00
191	LYS	CB	-40.57	-11.79	61.81	15.00
191	LYS	CG	-41.66	-10.76	61.63	15.00
191	LYS	CD	-42.98	-11.23	62.27	15.00
191	LYS	CE	-43.69	-12.29	61.41	15.00
191	LYS	NZ	-42.88	-13.54	61.16	15.00
191	LYS	С	-38.18	-12.50	61.33	15.00
191	LYS	0	-38.18	-13.07	62.44	15.00
192	GLY	N	-37.22	-12.67	60.43	15.00
192	GLY	CA	-36.06	-13.51	60.71	15.00
192	GLY	С	-36.27	-14.99	60.45	15.00
192	GLY	0	-35.41	-15.81	60.78	15.00
193	TYR		-37.40	-15.34	59.87	15.00
193	TYR	CA	-37.69	-16.74	59.58	15.00
193	TYR		-38.93	-17.20	60.33	15.00
193	TYR	CG	-38.68	-17.49	61.78	15.00
193	TYR		-38.81	-16.48	62.74	15.00
193	TYR		-38.56	-16.73	64.09	15.00
193	TYR		-38.31	-18.77	62.20	15.00
193	TYR		-38.06	-19.03	63.55	15.00
193	TYR		-38.18	-18.01	64.48	15.00
193	TYR		-37.91	-18.26	65.81	15.00
193	TYR		-37.92	-16.95	58.10	15.00
193	TYR		-38.22	-16.01	57.36	15.00
194	ILE		-37.81	-18.21	57.68	15.00
194	ILE		-38.04	-18.60	56.31	15.00
194	ILE		-36.73	-18.55	55.43	15.00
194	ILE		-35.60	-19.33	56.09	15.00
	ILE		-37.01	-19.10	54.03	15.00
194	ILE		-35.85	-18.96	53.07	15.00
194	ILE		-38.63	-20.00	56.30	15.00
194	ILE		-38.16	-20.88	57.01	15.00
195	LEU		-39.72	-20.18	55.55	15.00
195	LEU		-40.34	-21.49	55.42	15.00
195	LEU		-41.87	-21.37	55.28	15.00
195	LEU		-42.73	-20.47	56.17	15.00
195	LEU	CD1	-44.18	-20.60	55.73	15.00

195	LEU	CD2	-42.58	-20.81	57.65	15.00
195	LEU	С	-39.76	-22.05	54.13	15.00
195	LEU	0	-39.93	-21.45	53.06	15.00
196	MET	N	-39.03	-23.15	54.22	15.00
196	MET	CA	-38.43	~23.75	53.02	15.00
196	MET	CB	-36.94	-24.02	53.26	15.00
196	MET	CG	-36.14	-22.77	53.55	15.00
196	MET	SD	-34.44	-23.13	54.06	15.00
196	MET	CE	-33.70	-23.51	52.45	15.00
196	MET	С	-39.16	-25.05	52.64	15.00
196	MET	0	-39.62	-25.77	53.52	15.00
197	ALA	N	-39.26	-25.34	51.35	15.00
197	ALA	CA	-39.95	-26.53	50.86	15.00
197	ALA	CB	-39.82	-26.63	49.36	15.00
197	ALA	C	-39.52	-27.84	51.52	15.00
197	ALA	0	-38.32	-28.08	51.73	15.00
198	ARG	N	-40.50	-28.68	51.85	15.00
198	ARG	CA	-40.27	-29.98	52.48	15.00
198	ARG	CB	-41.04		53.81	15.00
198	ARG	CG	-41.06	-31.45	54.43	15.00
198	ARG	CD	-41.33	-31.38	55.94	15.00
198	ARG	NE	-42.61		56.26	15.00
	ARG		-43.75	-31.42	56.44	15.00
198	ARG	NH1	-43.78	-32.74	56.32	15.00
198	ARG	NH2	-44.87	-30.76	56.70	15.00
198	ARG	C	-40.73	-31.09	51.56	15.00
	ARG		-41.79	-31.00	50.96	15.00
	ASN		-39.95	-32.17	51.52	15.00
	ASN		-40.22	-33.33	50.68	15.00
	ASN		-41.58	-33.97	50.99	15.00
	ASN		-41.67	-34.52	52.39	15.00
	ASN		-40.69		53.14	15.00
	ASN		-42.85		52.77	15.00
	ASN		-40.16		49.21	15.00
	ASN				48.36	15.00
	LYS		-39.35		48.91	15.00
	LYS			-31.49	47.55	15.00
200			-39.47		47.43	15.00
	LYS			-29.64	46.22	15.00
	LYS		-41.12		46.46	15.00
	LYS			-28.18		15.00
	LYS			-28.02		
200	LYS	С	-37.78	-31.85	47.17	15.00

200 LYS O	-36.94	-30.98	46.93	15.00
201 ASN N	-37.51	-33.15	47.20	15.00
201 ASN CA	-36.21	-33.69	46.86	15.00
201 ASN CB	-35.94	-33.51	45.37	15.00
201 ASN CG	-36.89	-34.30	44.52	15.00
201 ASN OD1	-37.20	-35.46	44.81	15.00
201 ASN ND2	-37.40	-33.66	43.47	15.00
201 ASN C	-35.03	-33.16	47.67	15.00
201 ASN 0	-33.94	-32.96	47.11	15.00
202 ASN N	-35.23	-33.00	48.98	15.00
202 ASN CA	-34.18	-32.51	49.89	15.00
202 ASN CB	-33.04	-33.54	49.99	15.00
202 ASN CG	-32.05	-33.23	51.10	15.00
202 ASN OD1	-32.41	-32.67	52.14	15.00
202 ASN ND2	-30.79	-33.59	50.89	15.00
202 ASN C	-33.66	-31.16	49.41	15.00
202 ASN O	-32.46	-30.95	49.27	15.00
203 ALA N	-34.57	-30.23	49.18	15.00
203 ALA H	-35.42	-30.45	49.60	15.00
203 ALA CA	-34.31	-28.90	48.63	15.00
203 ALA CB	-35.55	-28.01	48.72	15.00
203 ALA C	-33.20	-28.21	49.44	15.00
203 ALA O	-33.27	-28.01	50.64	15.00
204 CYS N	-32.19	-27.72	48.68	15.00
204 CYS CA	-31.05	-26.98	49.22	15.00
204 CYS C	-30.21	-27.75	50.22	15.00
204 CYS 0	-29.44			15.00
204 CYS CB	-31.51		49.86	15.00
204 CYS SG	-32.47	-24.53	48.82	15.00
205 GLY N	-30.37	-29.07	50.27	15.00
205 GLY CA	-29.60	-29.89	51.20	15.00
205 GLY C	-30.01			15.00
205 GLY 0	-29.23		53.55	15.00
206 ILE N	-31.27		52.90	15.00
206 ILE CA		-29.38		
206 ILE CB		-29.05		
206 ILE CG2	-34.13		53.53	15.00
206 ILE CG1	-33.94		55.54	15.00
206 ILE CD1		-27.31	56.02	15.00
206 ILE C		-30.57		
206 ILE O			56.40	
207 ALA N		-31.77		
207 ALA CA	-31.27	-32.95	55.49	15.00

207	ALA	CB	_32 20	-33.99	55.31	15.00
	ALA		-29.89	-33.58	55.26	15.00
	ALA		-29.62	-34.70	55.70	15.00
	ASN		-28.99	-32.84	54.62	15.00
	ASN		-27.66	-33.36	54.34	15.00
	ASN		-27.05	-32.65	53.13	15.00
	ASN		-27.49	-33.27	51.83	15.00
	ASN		-27.92	-34.43	51.79	15.00
	ASN		-27.39	-32.51	50.75	15.00
	ASN		-26.67	-33.32	55.51	15.00
	ASN		-25.80	-34.19	55.61	15.00
	LEU		-26.82	-32.35	56.40	15.00
	LEU		-25.92	-32.20	57.53	15.00
209	LEU	CB	-24.79	-31.22	57.16	15.00
209	LEU	CG	-23.53	-31.18	58.02	15.00
209	LEU	CD1	-22.72	-32.44	57.77	15.00
209	LEU	CD2	-22.72	-29.93	57.68	15.00
209	LEU	С	-26.69	-31.68	58.75	15.00
209	LEU	0	-26.54	-30.52	59.15	15.00
210	ALA	N	-27.50	-32.55	59.34	15.00
210	ALA	H	-27.89	-32.98	58.56	15.00
210	ALA	CA	-28.32	-32.12	60.48	15.00
	ALA		-29.80	-32.37	60.21	15.00
	ALA		-27.94	-32.94	61.72	15.00
210	ALA	0	-27.57	-34.09	61.63	15.00
211		N	-28.10	-32.29	62.88	15.00
211			-27.80	-32.92	64.15	15.00
	SER		-26.28	-32.97	64.37	15.00
211			-25.71	-31.66	64.34	15.00
	SER		-28.46	-32.19	65.33	15.00
	SER		-29.01	-31.09	65.17	15.00
212			-28.43	-32.83	66.50	15.00
	PHE			-32.24	67.71	15.00
	PHE		-30.46	-32.59	67.86	15.00
	PHE			-34.06	67.93	15.00
	PHE			-34.79	66.77	15.00
	PHE			-34.72	69.16	15.00
212				-36.13	66.83	15.00
	PHE			-36.08 -36.79	69.22	15.00 15.00
	PHE			-36.79 -32.66	68.05	
212				-32.66 -33.77	68.96 69.03	15.00 15.00
212				-33.77	69.03 69.92	15.00
213	PRU	IA	-20.03	-31.76	ワフ・ブム	13.00

213	PRC	CD	-28.51	-30.37	69.98	15.00
213	PRC	CA	-27.28	-32.11	71.13	15.00
213	PRO	CB	-27.07	-30.76	71.79	15.00
213	PRO	CG	-28.33	-30.03	71.43	15.00
213	PRO	C	-28.10	-33.03	72.01	15.00
213	PRO	0	-29.33	-33.00	71.95	15.00
214	LYS	N	-27.42	-33.86	72.80	15.00
214	LYS	CA	-28.08	-34.78	73.73	15.00
214	LYS	CB	-27.64	-36.23	73.50	15.00
214	LYS	CG	-27.92	-36.75	72.10	15.00
214	LYS	CD	-27.72	-38.26	72.00	15.00
214	LYS	CE	-26.29	-38.66	72.30	15.00
214	LYS	NZ	-26.00	-39.99	71.69	15.00
214	LYS	C	-27.60	-34.34	75.10	15.00
214	LYS	0	-26.43	-34.00	75.26	15.00
215	MET	N	-28.50	-34.30	76.07	15.00
215	MET	CA	-28.12	-33.90	77.42	15.00
215	MET	CB	-28.97	-32.72	77.89	15.00
215	MET	CG	-28.96	-31.51	76.95	15.00
215	MET	SD	-29.63	-30.02	77.75	15.00
215	MET	CE	-28.68	-28.69	76.95	15.00
215	MET	С	-28.26	-35.09	78.36	15.00
215	MET	OT1	-27.93	-34.95	79.55	15.00
215	MET	OT2	-28.65	-36.17	77.89	15.00
216	HOH	OH2	-26.08	-16.55	83.97	15.00
217	HOH	OH2	-20.53	-32.33	79.43	15.00
218	HOH	OH2	-31.21	-16.22	65.49	15.00
219	HOH	OH2	-30.95	-18.19	68.23	15.00
220	нон	OH2	-6.96	-10.59	69.84	15.00
221	HOH	OH2	-15.23	-12.63	73.08	15.00
222	нон	OH2	-34.53	-23.51	69.96	15.00
223	НОН	OH2	-13.78	-33.08	69.63	15.00
224	НОН	OH2	-17.84	-17.71	5 <b>7</b> . 57	15.00
225	НОН	OH2	-24.92	-31.02	61.65	15.00
226	HOH	OH2	-12.76	-8.21	61.82	15.00
227	HOH	OH2	-14.16	-21.69	66.48	15.00
228	нон	OH2	-44.08	-26.87	48.48	15.00
229	нон	OH2	-44.49	-35.40	55.40	15.00
230	нон	OH2	-39.27	-16.80	68.54	15.00
231	нон	OH?	-24.12	-35.40	48.13	15.00
232	нон	OH2			63.42	
233	нон	OH2			44.36	15.00
234	нон	OH2	-27.99			

235 нон он2	-22.10	-30 02	61 67	15 00
236 HOH OH2	-27.35			15.00
237 нон он2	-29.19		71.93 70.74	15.00
238 HOH OH2	-29.55		83.52	15.00 15.00
239 НОН ОН2	-35.73		51.77	15.00
240 HOH OH2	-36.27		49.31	15.00
241 HOH OH2	-46.67		57.38	15.00
242 HOH OH2	-27.40			
243 HOH OH2		-15.97	60.76	15.00
244 HOH OH2	-18.00		62.85	15.00
245 HOH OH2	-33.49		70.56	15.00
246 HOH OH2	-44.87	-25.33	75.86	15.00
247 HOH OH2	-17.32		74.90	15.00
248 HOH OH2		-17.84	66.51	15.00
249 HOH OH2	-11.56	-21.89	82.27	15.00
250 НОН ОН2	-28.01	-35.21	58.24	15.00
251 HOH OH2	-35.05		53.00	15.00
252 нон он2	-31.64	-28.63	46.10	15.00
253 нон он2	-35.04	-24.79	46.85	15.00
254 нон он2	-41.38	-35.11	55.81	15.00
255 нон он2	-40.44	-19.77	71.52	15.00
256 нон он2	-43.66	-16.34	65.70	15.00
257 нон он2	-39.00	-11.99	70.39	15.00
258 нон он2	-30.92	-9.07	66.51	15.00
259 НОН ОН2	-32.51		60.41	15.00
260 НОН ОН2	-19.20	-8.29	62.91	15.00
261 HOH OH2	-33.67	-20.84	69.78	15.00
262 НОН ОН2	-32.87	-44.92	73.87	15.00
263 нон он2	-13.20	-24.01	76.81	
264 HOH OH2	-8.83		60.26	
265 HOH OH2	-17.23	-39.22	57.64	15.00
266 HOH OH2	-21.10	-32.62	61.09	15.00
267 HOH OH2		-33.44	60.85	15.00
268 HOH OH2	-6.37	-28.13	76.25	15.00
269 HOH OH2	-10.20	-38.51	65.40	15.00
270 HOH OH2 271 HOH OH2	-21.41 -22.56	-37.76 -39.95	78.27	15.00
271 HOH OH2 272 HOH OH2	-22.56 -30.18	-38.95 -25.13	69.33	15.00
272 HOR OH2 273 HOR OH2	-30.18 -12.08	-25.13 -12.20	93.77 63.63	15.00 15.00
274 HOH OH2	-1.36	-12.20 -9.62	67.96	15.00
275 HOH OH2	-28.39	-30.26	56.30	15.00
276 HOH OH2	-29.74	-20.19	48.42	15.00
277 HOH OH2	-26.12	-23.01	44.41	15.00
2 MOH OHZ		23.VI	22.24	13.00

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# TABLE IV 278 HOH OH2 -29.92 -34.21 47.42 15.00 279 HOH OH2 -26.24 -33.39 47.92 15.00 280 HOH OH2 -32.19 -28.29 42.42 15.00

281 HOH OH2 -37.49 -30.33 50.55 15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (Ų) for the cathepsin K complex with inhibitor (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide.

Residue	Atom	x	Y	Z	В
1 ALA	СВ	-44.33	-37.20	63.83	15.00
1 ALA	С	-46.76	-36.62	63.83	15.00
1 ALA	0	-47.46	-36.94	62.86	15.00
1 ALA	N	-46.07	-38.96	63.89	15.00
1 ALA	CA	-45.70	-37.59	64.36	15.00
2 PRO	N	-46.94	-35.47	64.51	15.00
2 PRO	CD	-46.25	-35.02	65.74	15.00
2 PRO	CA	-47.93	-34.49	64.07	15.00
2 PRO	CB	-47.63	-33.28	64.97	15.00
2 PRO	CG	-47.15	-33.90	66.23	15.00
2 PRO	С	-47.63	-34.15	62.63	15.00
2 PRO	0	-46.50	-34.30	62.16	15.00
3 ASP	N	-48.65	-33.74	61.88	15.00
3 ASP	CA	-48.39	-33.36	60.52	15.00
3 ASP	CB	-49.60	-33.60	59.63	15.00
3 ASP	CG	-49.78	-35.10	59.29	15.00
3 ASP		-50.65	-35.42	58.45	15.00
3 ASP	OD2	-49.05	-35.95	59.86	15.00
3 ASP	С	-47.92	-31.92	60.51	15.00
3 ASP	0	-47.44	-31.42	59.49	15.00
4 SER	N	-47.94	-31.30	61.69	15.00
4 SER	CA	-47.55	-29.91	61.87	15.00
4 SER	CB	-48.70	-28.98	61.49	15.00
4 SER	OG	-48.42	-27.64	61.85	15.00
4 SER	C	-47.13	-29.61	63.29	15.00
4 SER	0	-47.79	-30.03	64.24	15.00
5 VAL	N	-46.04	-28.86	63.45	15.00
5 VAL	CA	-45.60	-28.47	64.78	15.00
5 VAL	CB	-44.86	-29.61	65.53	15.00
5 VAL	CG1	-43.46	-29.81	64.97	15.00
5 VAL	CG2	-44.83	-29.31	67.02	15.00
5 VAL	C .	-44.78	-27.20	64.70	15.00
5 VAL	0	-44.00	-26.98	63.77	15.00

6 ASP N	-44.98	-26.34	65.69	15.00
6 ASP CA	-44.29	-25.07	65.76	15.00
6 ASP CB	-45.27	-23.94	65.39	15.00
6 ASP CG	-44.57	-22.65	65.00	15.00
6 ASP OD1	-43.38	-22.47	65.35	15.00
6 ASP OD2	-45.21	-21.81	64.33	15.00
6 ASP C	-43.72	-24.87	67.16	15.00
6 ASP O	-44.44	-24.51	68.10	15.00
7 TYR N	-42.41	-25.07	67.29	15.00
7 TYR CA	-41.75	-24.90	68.58	15.00
7 TYR CB	-40.35		68.57	15.00
7 TYR CG	-40.39	-27.00	68.75	15.00
7 TYR CD1	-40.49	-27.57	70.02	15.00
7 TYR CE1	-40.58	-28.94	70.20	15.00
7 TYR CD2	-40.38	-27.87	67.65	15.00
7 TYR CE2	-40.47	-29.25	67.81	15.00
7 TYR CZ	-40.57	-29.77	69.09	15.00
7 TYR OH	-40.68	-31.13	69.28	15.00
7 TYR C	-41.75	-23.46	69.08	15.00
7 TYR O	-41.62	-23.22	70.29	15.00
8 ARG N	-41.91	-22.51	68.17	15.00
8 ARG CA	-41.98	-21.10	68.55	15.00
8 ARG CB	-42.05	-20.19	67.33	15.00
8 ARG CG	-40.91	-20.36	66.38	15.00
8 ARG CD	-41.09	-19.45	65.19	15.00
8 ARG NE	-42.20	-19.81	64.32	15.00
8 ARG CZ	-42.56	-19.12	63.24	15.00
8 ARG NH1	-41.89	-18.03	62.89	15.00
8 ARG NH2	-43.63	-19.48	62.56	15.00
8 ARG C	-43.22	-20.92	69.42	15.00
8 ARG O	-43.17	-20.22	70.43	15.00
9 LYS N	-44.31	-21.59	69.04	15.00
9 LYS CA	-45.55	-21.53	69.81	15.00
9 LYS CB	-46.77	-21.98	68.99	15.00
9 LYS CG	-47.20	-21.00	67.88	15.00
9 LYS CD	-48.52	-21.46	67.22	15.00
9 LYS CE	-48.99	-20.55	66.08	15.00
9 LYS NZ	-49.38	-19.15	66.49	15.00
9 LYS C	-45.44	-22.32	71.13	15.00
9 LYS O	-46.27	-22.17	72.02	15.00
10 LYS N	-44.41	-23.16	71.23	15.00
10 LYS CA	-44.17	-23.94	72.43	15.00

10	LYS	СВ	-43.58	-25.32	72.11	15.00
10	LYS	CG	-44.58	-26.34	71.57	15.00
10	LYS	CD	-43.93	-27.72	71.47	15.00
10	LYS	CE	-44.97	-28.81	71.25	15.00
10	LYS	NZ	-45.93	-28.85	72.39	15.00
10	LYS	С	-43.25	-23.20	73.40	15.00
10	LYS	0	-43.06	-23.65	74.53	15.00
11	GLY	N	-42.67	-22.09	72.95	15.00
11	GLY	CA	-41.78	-21.32	73.79	15.00
11	GLY	С	-40.38	-21.91	73.87	15.00
11	GLY	0	-39.64	-21.67	74.83	15.00
12	TYR	N	-40.01	-22.66	72.84	15.00
12	TYR	CA	-38.71	-23.31	72.77	15.00
12	TYR	CB	-38.80	-24.65	72.05	15.00
12	TYR	CG	-39.27	-25.80	72.87	15.00
12	TYR	CD1	-40.41	-25.71	73.66	15.00
12	TYR	CE1	-40.84	-26.79	74.42	15.00
12	TYR	CD2	-38.57	-27.00	72.85	15.00
12	TYR	CE2	-38.99	-28.09	73.59	15.00
12	TYR	CZ	-40.12	-27.98	74.38	15.00
12	TYR	OH	-40.50	-29.07	75.14	15.00
12	TYR	С	-37.64	-22.48	72.08	15.00
12	TYR	0	-36.46	-22.81	72.16	15.00
13	VAL	N	-38.06	-21.44	71.37	15.00
13	VAL	CA	-37.14	-20.62	70.58	15.00
13	VAL	CB	-37.66	-20.50	69.13	15.00
13	VAL	CG1	-36.66	-19.77	68.25	15.00
13	VAL	CG2	-37.95	-21.87	68.56	15.00
13	VAL		-36.89	-19.23	71.14	15.00
13	VAL	0	-37.84	-18.50	71.45	15.00
14	THR	N	-35.63	-18.84	71.23	15.00
	THR		-35.31	-17.51	71.72	15.00
14	THR	CB	-33.91	-17.47	72.36	15.00
14	THR	OG1	-32.93	-17.77	71.36	15.00
14	THR	CG2	-33.80	-18.49	73.47	15.00
14	THR	C	-35.44	-16.50	70.56	15.00
14	THR	0	-35.63	-16.89	69.40	15.00
15	PRO	N	-35.41	-15.19	70.86	15.00
15	PRO	CD	-35.34	-14.54	72.18	15.00
15	PRO	CA	-35.53	-14.18	69.79	15.00
15	PRO	СВ	-35.29	-12.87	70.53	15.00
15	PRO	CG	-35.87	-13.15	71.89	15.00

15 PRO C	-34.52	-14.37	68.66	15.00
15 PRO O	-33.50	-15.04	68.82	15.00
16 VAL N	-34.82	-13.76	67.52	15.00
16 VAL CA	-33.93	-13.83	66.37	15.00
16 VAL CB	-34.62	-13.31	65.09	15.00
16 VAL CG1	-33.61	-13.11	63.98	15.00
16 VAL CG2	-35.68	-14.31	64.65	15.00
16 VAL C	-32.68	-13.01	66.64	15.00
16 VAL O	-32.76	-11.88	67.11	15.00
17 LYS N	-31.52	-13.61	66.39	15.00
17 LYS CA	-30.24	-12.96	66.58	15.00
17 LYS CB	-29.25	-13.89	67.30	15.00
17 LYS CG	-29.81	-14.64	68.50	15.00
17 LYS CD	-30.24	-13.71	69.61	15.00
17 LYS CE	-30.58	-14.46	70.88	15.00
17 LYS NZ	-31.75	-15.34	70.73	15.00
17 LYS C	-29.67	-12.53	65.23	15.00
17 LYS O	-30.20	-12.88	64.17	15.00
18 ASN N	-28.57	-11.79	65.27	15.00
18 ASN CA	-27.90	-11.32	64.06	15.00
18 ASN CB	-28.17	-9.84	63.80	15.00
18 ASN CG	-27.66		62.45	15.00
18 ASN OD1	-26.79	-10.02	61.85	15.00
18 ASN ND2	-28.20	-8.29	61.95	15.00
18 ASN C	-26.41	-11.58	64.19	15.00
18 ASN 0	-25.74	-11.03	65.08	15.00
19 GLN N	-25.89	-12.42	63.30	15.00
19 GLN CA	-24.48	-12.75	63.31	15.00
19 GLN CB	-24.20	-14.02	62.48	15.00
19 GLN CG	-24.56	-13.94	61.00	15.00
19 GLN CD	-24.28	-15.24	60.27	15.00
19 GLN OE1		-15.79	59.60	15.00
19 GLN NE2	-23.06	-15.74	60.40	15.00
19 GLN C		-11.60	62.86	15.00
19 GLN 0		-11.51	63.27	15.00
20 GLY N	-24.12	-10.71	62.03	15.00
20 GLY CA	-23.32	-9.59	61.55	15.00
20 GLY C	-22.33	-10.05	60.49	15.00
20 GLY 0	-22.59	-11.03	59.78	15.00
21 GLN N	-21.19	-9.38		15.00
21 GLN CA	-20.18	-9.73	59.39	15.00
21 GLN CB	-19.51	-8.48	58.81	15.00

21	GLN	CG	-20.42	-7.63	57.93	15.00
21	GLN	CD	-20.79	-8.33	56.62	15.00
21	GLN	OE1	-20.02	-9.12	56.07	15.00
21	GLN	NE2	-21.97	-8.02	56.11	15.00
21	GLN	С	-19.15	-10.71	59.95	15.00
21	GLN	0	-17.96	-10.40	60.02	15.00
22	CYS	N	-19.63	-11.88	60.34	15.00
22	CYS	CA	-18.79	-12.94	60.89	15.00
22	CYS	С	-19.49	-14.24	60.53	15.00
22	CYS	0	-20.71	-14.34	60.62	15.00
22	CYS	CB	-18.61	-12.78	62.41	15.00
22	CYS	SG	-18.03	-14.24	63.33	15.00
23	GLY	N	-18.73	-15.19	60.00	15.00
23	GLY	CA	-19.29	-16.48	59.66	15.00
23	GLY	С	-19.53	-17.35	60.89	15.00
23	GLY	0	-19.04	-18.48	60.98	15.00
24	SER	N	-20.36	-16.86	61.81	15.00
24	SER	CA	-20.67	-17.60	63.03	15.00
24	SER	CB	-20.61	-16.66	64.22	15.00
24	SER	OG	-21.35	-15.49	63.95	15.00
24	SER	С	-22.01	-18.36	62.96	15.00
24	SER	0	-22.58	-18.73	63.99	15.00
25	CYS	N	-22.50	-18.58	61.74	15.00
25	CYS	CA	-23.76	-19.28	61.52	15.00
25	CYS	CB	-23.98	-19.51	60.01	15.00
25	CYS	SG	-22.57	-20.30	59.15	15.00
25	CYS	С	-23.86	-20.58	62.32	15.00
25	CYS	0	-24.84	-20.82	63.02	15.00
25	INH	C1	-28.50	-9.52	55.70	15.00
25	INH	C2	-28.63	-9.33	57.08	15.00
25	INH	C3	-27.56	-9.63	57.94	15.00
25	INH	C4	-26.35	-10.11	57.43	15.00
25	INH	C5	-26.23	-10.29	56.05	15.00
25	INH	C6	-27.29	-10.00	55.19	15.00
25	INH	C7	-25.20	-10.40	58.35	15.00
25	INH	08	-24.73	-11.77	58.36	15.00
25	INH	C9	-24.03	-12.42	57.30	15.00
25	INH	010	-24.33	-12.23	56.10	15.00
25	INH	C11	-22.27	-14.01	56.70	15.00
25	INH	C12	-20.77	-13.63	56.80	15.00
25	INH	C13	-20.18	-12.60	55.82	15.00
25	INH	C14	-19.01	-11.83	56.47	15.00

25		E C15	-21.22	-11.63	55.23	15.00
25		I C16	-22.50	-15.59	56.80	15.00
25		I S17	-23.78	-16.32	55.92	15.00
25		1 N18	-21.80	-16.55	57.50	15.00
25		I C19	-22.21	-17.87	57.39	15.00
25	INE	N20	-23.05	-13.25	57.68	15.00
25	INH	C21	-23.27	-17.88	56.55	15.00
25	INH	C22	-21.58	-19.10	58.24	15.00
25	INH	023	-21.37	-18.39	59.17	15.00
25	INH	C24	-13.79	-23.51	54.96	15.00
25	INH	C25	-14.23	-22.84	56.08	15.00
25	INH	C26	-14.83	-23.54	57.12	15.00
25	INH	C27	-15.00	-24.93	57.04	15.00
25	INH	C28	-14.54	-25.60	55.91	15.00
25	INH	C29	-13.94	-24.90	54.87	15.00
25	INH	C30	-15.72	-25.67	58.14	15.00
25	INH	031	-17.10	-25.93	57.71	15.00
25	INH	C32	-17.91	-25.03	56.96	15.00
25	INH	033	-17.69	-24.81	55.77	15.00
25	INH	C34	-19.82	-23.49	57.00	15.00
25		C35	-21.22	-24.12	56.84	15.00
25		C36	-21.92	-24.89	57.97	15.00
25		C37	-21.43	-26.31	58.12	15.00
25		C38	-21.86	-24.15	59.29	15.00
25	INH	C39	-19.87	-22.15	57.76	15.00
25		040	-19.60	-22.13	58.96	15.00
25		N41	-20.18	-21.00	57.08	15.00
25	INH	N42	-20.20	-19.65	57.78	15.00
25	INH	N43	-18.90	-24.44	57.63	15.00
	TRP	N	-22.80	-21.38	62.25	15.00
26	TRP	CA	-22.73	-22.65	62.97	15.00
26		CB	-21.39	-23.33	62.67	15.00
26	TRP		-20.19	-22.46	62.98	15.00
	TRP		-19.41	-22.45	64.19	15.00
		CE2	-18.44	-21.44	64.05	15.00
	TRP	CE3	-19.43	-23.21	65.37	15.00
26	TRP	CD1	-19.67	-21.48	62.19	15.00
26	TRP	NE1	-18.62	-20.86	62.82	15.00
	TRP	CZ2	-17.50	-21.15	65.06	15.00
26	TRP	CZ3	-18.50	-22.92	66.37	15.00
	TRP	CH2	-17.55	-21.91	66.21	15.00
26	TRP	C	-22.95	-22.50	64.49	15.00

26	TRP	0	-23.65	-23.30	65.10	15.00
27	ALA	N	-22.35	-21.46	65.08	15.00
27	ALA	CA	-22.45	-21.18	66.51	15.00
27	ALA	CB	-21.56	-20.02	66.89	15.00
27	ALA	С	-23.90	-20.90	66.88	15.00
27	ALA	0	-24.46	-21.56	67.74	15.00
28	PHE	N	-24.53	-19.94	66.19	15.00
28	PHE	CA	-25.93	-19.58	66.41	15.00
28	PHE	CB	-26.36	-18.44	65.48	15.00
28	PHE	CG	-25.83	-17.09	65.88	15.00
28	PHE	CD1	-24.66	-16.59	65.32	15.00
28	PHE	CD2	-26.48	-16.32	66.84	15.00
28	PHE	CE1	-24.14	~15.37	65.71	15.00
28	PHE	CE2	-25.97	-15.09	67.24	15.00
28	PHE	CZ	-24.80	-14.62	66.67	15.00
28	PHE	С	-26.87	-20.78	66.23	15.00
28	PHE	0	-27.85	-20.92	66.96	15.00
29	SER	N	-26.56	-21.64	65.26	15.00
29	SER	CA	-27.35	-22.83	64.99	15.00
29	SER	CB	-26.95	-23.45	63.64	15.00
29	SER	OĞ	-27.68	-24.63	63.36	15.00
29	SER	С	-27.21	-23.84	66.13	15.00
29	SER	0	-28.19	-24.49	66.50	15.00
30	SER	N	-26.01	-23.96	66.69	15.00
30	SER	CA	-25.73	-24.88	67.79	15.00
30	SER	CB	-24.23	-25.10	68.00	15.00
30	SER	OG	-23.60	-25.57	66.82	15.00
30	SER	С	-26.37	-24.39	69.09	15.00
30	SER	0	-26.95	-25.16	69.85	15.00
31	VAL	N	-26.23	-23.09	69.33	15.00
31	VAL	CA	-26.80	-22.45	70.51	15.00
31	VAL		-26.39	-20.95	70.55	15.00
	VAL		-27.42	-20.12	71.24	15.00
31	VAL	CG2	-25.07	-20.81	71.26	15.00
	VAL		-28.31	-22.63	70.48	15.00
31	VAL	0	-28.91	-22.92	71.51	15.00
32	GLY	N	-28.89	-22.52	69.29	15.00
32	GLY	CA	-30.32	-22.68	69.11	15.00
32	GLY	С	-30.78	-24.09	69.45	15.00
32	GLY	0	-31.86	-24.28	69.99	15.00
33	ALA	N	-29.97	-25.08	69.09	15.00
33	ALA	CA	-30.29	-26.47	69.38	15.00

33	ALA	CB	-29.33	-27.39	68.65	15.00
33	ALA	C	-30.19		70.89	15.00
33	ALA	. 0	~31.08	-27.30	71.50	15.00
34	LEU	N	-29.13	-26.18	71.50	15.00
34	LEU	CA	-28.93	-26.32	72.93	15.00
34	LEU	CB	-27.58	-25.75	73.35	15.00
34	LEU	CG	-26.31	-26.50	72.95	15.00
34	LEU	CD1	-25.08	-25.64	73.21	15.00
34	LEU	CD2	-26.24	-27.79	73.73	15.00
34	LEU	C	-30.07	-25.65	73.69	15.00
34	LEU	0	-30.59	-26.21	74.64	15.00
35	GLU	N	-30.47	-24.47	73.22	15.00
35	GLU	CA	-31.55	-23.70	73.82	15.00
35	GLU	CB	-31.77	-22.39	73.08	15.00
35	GLU	CG	-30.84	-21.28	73.54	15.00
35	GLU	CD	-30.76	-20.13	72.55	15.00
35	GLU	OE1	-31.51	-20.14	71.54	15.00
35	GLU	OE2	-29.93	-19.22	72.76	15.00
35	GLU	С	-32.86	-24.47	73.95	15.00
35	GLU		-33.52	-24.39	75.00	15.00
36	GLY	N	-33.21	-25.21	72.90	15.00
	GLY	CA	-34.42	-26.00	72.90	15.00
	GLY	C	-34.35	-27.13	73.91	15.00
36	GLY		-35.29	-27.37	74.66	15.00
37	GLN		-33.22	-27.82	73.95	15.00
37	GLN	CA	-33.04	-28.92	74.90	15.00
37	GLN		-31.77	-29.71	74.56	15.00
37	GLN		-31.84	-30.38	73.19	15.00
37	GLN		-33.17	-31.11	72.97	15.00
37	GLN		-33.60	-31.90	73.81	15.00
37	GLN	NE2	-33.82	-30.83	71.85	15.00
37	GLN		-33.05	-28.41	76.35	15.00
37	GLN		-33.63	-29.04	77.23	15.00
38	LEU		-32.45	-27.24	76.57	15.00
38			-32.42	-26.63	77.90	15.00
38	LEU	CB	-31.61	-25.34	77.89	15.00
38	LEU		-31.50	-24.54	79.20	15.00
38	LEU			-25.41	80.34	15.00
38	LEU			-23.34	78.95	15.00
38	LEU			-26.35	78.35	15.00
38	LEU			-26.60	79.50	15.00
39	LYS	N	-34.66	-25.84	77.42	15.00

39	LYS	CA	-36.06	-25.56	77.68	15.00
39	LYS	CB	-36.71	-24.85	76.49	15.00
39	LYS	CG	-38.21	-24.62	76.62	15.00
39	LYS	CD	-38.52	-23.64	77.74	15.00
39	LYS	CE	-40.03	-23.47	77.92	15.00
39	LYS	NZ	-40.35	-22.39	78.91	15.00
39	LYS	С	-36.83	-26.83	78.04	15.00
39	LYS	0	-37.55	-26.89	79.04	15.00
40	LYS	N	-36.65	-27.87	77.23	15.00
40	LYS	CA	-37.33	-29.14	77.44	15.00
40	LYS	CB	-37.06	-30.09	76.28	15.00
40	LYS	CG	-37.54	-31.50	76.53	15.00
40	LYS	CD	-37.53	-32.32	75.26	15.00
40	LYS	CE	-38.47	-31.72	74.22	15.00
40	LYS	NZ	-38.75	-32.66	73.09	15.00
40	LYS	C	-37.01	-29.80	78.78	15.00
40	LYS	0	-37.92	-30.20	79.52	15.00
41	LYS	N	-35.73	-29.90	79.11	15.00
41	LYS	CA	-35.29	-30.52	80.36	15.00
41	LYS	CB	-33.84	-31.02	80.22	15.00
41	LYS	CG	-33.70	-32.50	79.87	15.00
41	LYS	CD	-34.49	-32.90	78.62	15.00
41	LYS	CE	-33.58	-33.12	77.41	15.00
41	LYS	NZ	-33.06	-31.85	76.83	15.00
41	LYS	С	-35.47	-29.73	81.68	15.00
41	LYS	0	-35.74	-30.32	82.73	15.00
42	THR		-35.28	-28.40	81.63	15.00
42	THR		-35.41	-27.57	82.83	15.00
	THR		-34.17	-26.65	83.04	15.00
42	THR		-34.20	-25.57	82.10	15.00
42	THR		-32.87	-27.43	82.84	15.00
42	THR		-36.64	-26.66	82.82	15.00
	THR		-37.07	-26.16	83.86	15.00
	GLY			-26.41	81.63	15.00
	GLY		-38.33	-25.55	81.52	15.00
	GLY		-37.93	-24.10	81.41	15.00
	GLY		-38.78	-23.23	81.26	15.00
	LYS		-36.63	-23.82	81.53	15.00
	LYS		-36.11	-22.46	81.41	15.00
	LYS		-34.91		82.33	15.00
	LYS		-35.25		83.79	15.00
44	LYS	CD	-34.06	-21.63	84.59	15.00

44 LYS CE	-33.69	-20.18	84.17	15.00
44 LYS NZ	-32.77	-19.50	85.16	15.00
44 LYS C	-35.72	-22.17	79.96	15.00
<b>44</b> LYS 0	-35.24	-23.05	79.25	15.00
45 LEU N	-35.91	-20.93	79.54	15.00
45 LEU CA	-35.56	-20.52	78.19	15.00
45 LEU CB	-36.80	-20.28	77.31	15.00
45 LEU CG	-36.49	-19.86	75.87	15.00
45 LEU CD1	-36.01	-21.07	75.08	15.00
45 LEU CD2	-37.69	-19.23	75.17	15.00
45 LEU C	-34.71	-19.26	78.25	15.00
45 LEU O	-35.22	-18.17	78.53	15.00
46 LEU N	-33.41	-19.42	78.02	15.00
46 LEU CA	-32.50	-18.29	78.00	15.00
46 LEU CB	-31.75	-18.14	79.35	15.00
46 LEU CG	-31.05	-19.27	80.11	15.00
46 LEU CD1	-32.04	-20.12	80.86	15.00
46 LEU CD2	-30.23	-20.09	79.17	15.00
46 LEU C	-31.54	-18.34	76.80	15.00
46 LEU O	-31.41	-19.37	76.14	15.00
47 ASN N	-30.93	-17.20	76.48	15.00
47 ASN CA	-30.00	-17.12	75.36	15.00
47 ASN CB	-29.88	-15.69	74.84	15.00
47 ASN CG	-31.21	-15.12	74.46	15.00
47 ASN OD1	-31.91	-15.66	73.60	15.00
47 ASN ND2	-31.59	-14.04	75.11	15.00
47 ASN C	-28.64	-17.64	75.75	15.00
47 ASN O	-28.10	-17.24	76.78	15.00
48 LEU N	-28.12	-18.56	74.94	15.00
48 LEU CA	-26.82	-19.15	75.18	15.00
48 LEU CB	-26.80	-20.63	74.80	15.00
48 LEU CG	-27.73	-21.45	75.71	15.00
48 LEU CD1	-27.55	-22.93	75.44	15.00
48 LEU CD2	-27.44	-21.15	77.17	15.00
48 LEU C	-25.73	-18.36	74.47	15.00
48 LEU O	-26.04	-17.44	73.71	15.00
49 SER N	-24.48	-18.72	74.69	15.00
49 SER CA	-23.36	-18.00	74.12	15.00
49 SER CB	-22.27	-17.78	75.18	15.00
49 SER OG	-21.19	-17.01	74.69	15.00
49 SER C	-22.73	-18.50	72.83	15.00
49 SER 0	-21.93	-19.43	72.84	15.00

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50	PRO	N	-23.07	-17.87	71.68	15.00
50	PRO	CD	-24.14	-16.88	71.44	15.00
50	PRO	CA	-22.47	-18.30	70.42	15.00
50	PRO	CB	-23.33	-17.58	69.36	15.00
50	PRO	CG	-23.81	-16.35	70.07	15.00
50	PRO	С	-21.00	-17.83	70.39	15.00
50	PRO	0	-20.15	-18.48	69.77	15.00
51	GLN	N	-20.70	-16.73	71.10	15.00
51	GLN	CA	-19.34	-16.18	71.19	15.00
51	GLN	CB	-19.35	-14.78	71.82	15.00
51	GLN	CG	-18.06	-13.96	71.66	15.00
51	GLN	CD	-17.76	-13.49	70.22	15.00
51	GLN	OE1	-18.60	-12.89	69.54	15.00
51	GLN	NE2	-16.54	-13.74	69.77	15.00
51	GLN	С	-18.42	-17.16	71.95	15.00
51	GLN	0	-17.25	-17.33	71.59	15.00
52	asn	N	-18.98	-17.86	72.94	15.00
52	ASN	CA	-18.23	-18.85	73.70	15.00
52	asn	CB	-19.14	-19.54	74.73	15.00
52	ASN	CG	-18.40	-20.50	75.66	15.00
52	ASN	OD1	-18.99	-21.02	76.61	15.00
52	ASN	ND2	-17.11	-20.73	75.41	15.00
52	ASN	С	-17.68	-19.86	72.70	15.00
52	ASN	0	-16.50	-20.21	72.73	15.00
53	LEU	N	-18.55	-20.28	71.78	15.00
53	LEU	CA	-18.19	-21.24	70.74	15.00
53	LEU	CB	-19.44	-21.79	70.06	15.00
53	LEU	CG	-20.11	-23.02	70.67	15.00
53	LEU	CD1	-20.05	-22.96	72.19	15.00
53	LEU	CD2	-21.55	-23.11	70.17	15.00
53	LEU	С	-17.21	-20.66	69.72	15.00
53	LEU	0	-16.19	-21.29	69.42	15.00
54	VAL	N	-17.51	-19.46	69.21	15.00
54	VAL	CA	-16.65	-18.82	68.22	15.00
54	VAL	CB	-17.16	-17.39	67.85	15.00
54	VAL	CG1	-16.22	-16.73	66.85	15.00
54	VAL	CG2	-18.56	-17.47	67.25	15.00
54	VAL	С	-15.19	-18.75	68.68	15.00
54	VAL	0	-14.28	-19.13	67.94	15.00
55	ASP	N	-14.99	-18.33	69.93	15.00
55	ASP	CA	-13.65	-18.18	70.52	15.00
55	ASP	СВ	-13.64	-17.12	71.64	15.00

55 ASP CG	-14.11	-15.73	71.18	15.00
55 ASP OD1	-14.12	-15.43	69.96	15.00
55 ASP OD2	-14.46	-14.93	72.06	15.00
55 ASP C	-12.98	-19.44	71.08	15.00
55 ASP 0	-11.75	-19.53	71.15	15.00
56 CYS N	-13.79	-20.43	71.47	15.00
56 CYS CA	-13.26	-21.63	72.12	15.00
56 CYS C	-13.13	-22.97	71.41	15.00
56 CYS O	-12.36	-23.82	71.86	15.00
56 CYS CB	-13.97	-21.81	73.45	15.00
56 CYS SG	-13.91	-20.34	74.55	15.00
57 VAL N	-13.92	-23.20	70.36	15.00
57 VAL CA	-13.85	-24.48	69.64	15.00
57 VAL CB	-15.13	-24.77	68.83	15.00
57 VAL CG1	-15.08	-26.20	68.30	15.00
57 VAL CG2	-16.37	-24.52	69.66	15.00
57 VAL C	-12.67	-24.45	68.68	15.00
57 VAL O	-12.73	-23.82	67.62	15.00
58 SER N	-11.60	-25.15	69.04	15.00
58 SER CA	-10.40	-25.18	68.22	15.00
58 SER CB	-9.19	-25.66	69.02	15.00
58 SER OG	-9.56	-26.66	69.95	15.00
58 SER C	-10.54	-25.93	66.91	15.00
58 SER O	-9.71	-25.75	66.02	15.00
59 GLU N	-11.56	-26.78	66.79	15.00
59 GLU CA	-11.79	-27.55	65.57	15.00
59 GLU CB	-12.53	-28.86	65.84	15.00
59 GLU CG	-11.72	-29.95	66.56	15.00
59 GLU CD	-11.47	-29.63	68.03	15.00
59 GLU OE1	-12.44	-29.48	68.79	15.00
59 GLU OE2	-10.28	-29.54	68.42	15.00
59 GLU C	-12.51	-26.74	64.50	15.00
59 GLU O	-12.45	-27.06	63.32	15.00
60 ASN N	-13.22	-25.69	64.92	15.00
60 ASN CA	-13.91	-24.83	63.98	15.00
60 ASN CB	-15.29	-24.45	64.49	15.00
60 ASN CG	-16.25	-25.62	64.51	15.00
60 ASN OD1	-17.17	-25.66	65.32	15.00
60 ASN ND2	-16.04	-26.59	63.62	15.00
60 ASN C	-13.03	-23.63	63.72	15.00
60 ASN O	-12.01	-23.46	64.39	15.00
61 ASP N	-13.39	-22.81	62.74	15.00

61	ASP (	CA	-12.56	-21.66	62.39	15.00
61	ASP (	CB ·	-12.27	-21.64	60.88	15.00
61	ASP (	CG ·	-11.96	-23.05	60.30	15.00
61	ASP (	) I O	-12.89	-23.70	59.76	15.00
61	ASP (	DD2 -	-10.78	-23.50	60.37	15.00
61	ASP (		-13.12	-20.30	62.86	15.00
61	ASP (	,	-12.75	-19.26	62.32	15.00
62	GLY N	1 -	-13.97	-20.31	63.88	15.00
62	GLY C	:A -	-14.54	-19.06	64.36	15.00
62	GLY C	: -	-15.40	-18.39	63.30	15.00
62	GLY C	) -	-16.41	-18.95	62.84	15.00
63	CYS N	<b>.</b>	-15.00	-17.18	62.90	15.00
63	CYS C	:A -	-15.71	-16.43	61.88	15.00
63	CYS C	: -	15.44	-16.96	60.47	15.00
63	CYS C	) -	16.00	-16.47	59.49	15.00
63	CYS C	:B -	15.39	-14.94	61.97	15.00
63	CYS S	:G -	16.00	-14.06	63.45	15.00
64	GLY N	1 -	14.57	-17.97	60.38	15.00
64	GLY C	:A -	14.27	-18.57	59.09	15.00
64	GLY C	: -	15.01	-19.88	58.84	15.00
64	GLY C	) -	14.59	-20.68	58.01	15.00
65	GLY N	r -	16.09	-20.12	59.59	15.00
65	GLY C			-21.34	59.42	15.00
65	GLY C	: -	16.66	-22.39	60.49	15.00
65	GLY O	•	15.77	-22.30	61.34	15.00
66	GLY N	-	17.52	-23.40	60.46	15.00
66	GLY C			-24.49	61.42	15.00
66	GLY C	: -	18.61	-25.44	61.32	15.00
66	GLY O		19.49	-25.27	60.48	15.00
67	TYR N	-	18.62	-26.44	62.18	15.00
67	TYR C			-27.44	62.21	15.00
67	TYR C			-28.83	61.87	15.00
67	TYR C		18.68	-28.95	60.44	15.00
	TYR C		19.58	-29.32	59.44	15.00
	TYR C			<b>-29.3</b> 9	58.11	15.00
67	TYR C			-28.64	60.07	15.00
67	TYR C			-28.70	58.75	15.00
67	TYR C			-29.07	57.77	15.00
	TYR O			-29.10	56.45	15.00
67	TYR C			-27.48	63.5€	15.00
67				-27.20	64.59	15.00
83	MET N	-	21.65	-27.77	63.56	15.00

68 MET CA	-22.44	-27.85	64.78	15.00
68 MET CB	-23.93	-28.03	64.44	15.00
68 MET CG	-24.58	-26.86	63.66	15.00
68 MET SD	-24.01	-26.58	61.93	15.00
68 MET CE	-25.23	-27.50	61.01	15.00
68 MET C	-21.91	-28.98	65.68	15.00
68 MET O	-21.68	-28.77	66.87	15.00
69 THR N	-21.68	-30.16	65.08	15.00
69 THR CA	-21.17	-31.32	65.81	15.00
69 THR CB	-20.89	-32.54	64.87	15.00
69 THR OG1	-20.07	-32.14	63.76	15.00
69 THR CG2	-22.20	-33.12	64.35	15.00
69 THR C	-19.91	-31.01	66.64	15.00
69 THR O	-19.80	-31.46	67.78	15.00
70 ASN N	-18.97	-30.25	66.06	15.00
70 ASN CA	-17.74	-29.87	66.77	15.00
70 ASN CB	-16.76	-29.07	65.89	15.00
70 ASN CG	-16.17	-29.89	64.75	15.00
70 ASN OD1	-15.66	-29.34	63.79	15.00
70 ASN ND2	-16.25	-31.20	64.86	15.00
70 ASN C	-18.11	-29.01	67.98	15.00
70 ASN 0	-17.57	-29.19	69.08	15.00
71 ALA N	-19.02	-28.07	67.74	15.00
71 ALA CA	-19.50	-27.16	68.77	15.00
71 ALA CB	-20.44	-26.11	68.16	15.00
71 ALA C	-20.20	-27.92	69.91	15.00
71 ALA O	-20.03	-27.59	71.08	15.00
72 PHE N	-20.95	-28.97	69.56	15.00
72 PHE CA	-21.63	-29.82	70.54	15.00
72 PHE CB	-22.65	-30.75	69.86	15.00
72 PHE CG	-23.80	-30.02	69.25	15.00
72 PHE CD1	-24.32	-28.88	69.86	15.00
72 PHE CD2	-24.37	-30.48	68.08	15.00
72 PHE CE1	-25.40	-28.21	69.30	15.00
72 PHE CE2	-25.46	-29.81	67.51	15.00
72 PHE CZ	-25.97	-28.67	68.13	15.00
72 PHE C	-20.64	-30.63	71.36	15.00
72 PHE 0	-20.69	-30.66	72.59	15.00
73 GLN N	-19.71	-31.27	70.67	15.00
73 GLN CA	-18.70	-32.08	71.33	15.00
73 GLN CB	-17.83	-32.79	70.31	15.00
73 GLN CG	-16.90	-33.82	70.92	15.00

73	GLN	CD	-16.28	-34.73	69.87	15.00
73	GLN	OE1	-16.83	-34.92	68.78	15.00
73	GLN	NE2	-15.14	-35.30	70.20	15.00
73	GLN	С	-17.87	-31.21	72.25	15.00
73	GLN	0	-17.49	-31.64	73.33	15.00
74	TYR	N	-17.60	-29.98	71.82	15.00
74	TYR	CA	-16.83	-29.03	72.62	15.00
74	TYR	CB	-16.61	-27.69	71.89	15.00
74	TYR	CG	-16.39	-26.51	72.83	15.00
74	TYR	CD1	-15.18	-26.35	73.51	15.00
74	TYR	CE1	-15.02	-25.35	74.47	15.00
74	TYR	CD2	-17.43	-25.63	73.12	15.00
74	TYR	CE2	-17.28	-24.64	74.08	15.00
74	TYR	CZ	-16.08	-24.50	74.75	15.00
74	TYR	OH	-15.96	-23.57	75.75	15.00
74	TYR	С	-17.55	-28.79	73.94	15.00
74	TYR	0	-16.92	-28.75	75.00	15.00
75	VAL	N	-18.87	-28.62	73.86	15.00
75	VAL	CA	-19.70	-28.36	75.04	15.00
75	VAL	CB	-21.11	-27.88	74.63	15.00
	VAL		-21.96	-27.66	75.85	15.00
	VAL		-21.00	-26.57	73.83	15.00
75	VAL	С	-19.77	-29.55	75.98	15.00
75	VAL	0	-19.91	-29.38	77.19	15.00
76	GLN		-19.66	-30.74	75.42	15.00
	GLN		-19.66	-31.95	76.22	15.00
	GLN		-19.84	-33.21	75.36	15.00
	GLN		-19.80	-34.51	76.14	15.00
	GLN		-19.78	-35.75	75.25	15.00
76	GLN		-19.34	-35.70	74.09	15.00
	GLN		-20.24	-36.86	75.79	15.00
	GLN		-18.34	-32.00	77.00	15.00
	GLN		-18.34	-32.09	78.22	15.00
77	LYS		-17.22	-31.90	76.29	15.00
77	LYS		-15.89	-31.95	76.91	15.00
77	LYS		-14.79	-32.01	75.85	15.00
77	LYS		-14.77	-33.29	75.01	15.00
77	LYS		-13.80	-33.14	73.84	15.00
77	LYS		-13.62	-34.44	73.09	15.00
77			-12.55	-34.33	72.06	15.00
77			-15.63	-30.80	77.87	15.00
77	LYS	0	-14.99	-30.98	78.91	15.00

78 ASN N	-16.09	-29.61	77.50	15.00
78 ASN CA	-15.91	-28.43	78.32	15.00
78 ASN CB	-16.18	-27.16	77.51	15.00
78 ASN CG	-15.95	-25.89	78.31	15.00
78 ASN OD1	-14.90		78.92	15.00
78 ASN ND2	-16.93	-24.99	78.28	15.00
78 ASN C	-16.83	-28.52	79.52	15.00
78 ASN O	-16.64	-27.81	80.49	15.00
79 ARG N	-17.81	-29.42	79.44	15.00
79 ARG CA	-18.82	-29.64	80.47	15.00
79 ARG CB	-18.19	-30.06	81.81	15.00
79 ARG CG	-17.69	-31.48	81.85	15.00
79 ARG CD	-16.75	-31.70	83.00	15.00
79 ARG NE	-16.07	-32.98	82.88	15.00
79 ARG CZ	-14.79	-33.13	82.56	15.00
79 ARG NH1	-14.02	-32.08	82.33	15.00
79 ARG NH2	-14.28	-34.36	82.41	15.00
79 ARG C	-19.77	-28.45	80.65	15.00
79 ARG O	-20.43	-28.32	81.69	15.00
80 GLY N	-19.84	-27.58	79.66	15.00
80 GLY CA	-20.72	-26.45	79.77	15.00
80 GLY C	-20.50	-25.38	78.73	15.00
80 GLY O	-19.60	-25.49	77.89	15.00
81 ILE N	-21.38	-24.39	78.75	15.00
81 ILE CA	-21.34	-23.24	77.85	15.00
81 ILE CB	-22.12	-23.47	76.52	15.00
81 ILE CG2	-23.54	-24.00	76.80	15.00
81 ILE CG1	-22.15	-22.16	75.71	15.00
81 ILE CD1	-22.81	-22.28	74.36	15.00
81 ILE C	-21.98	-22.09	78.60	15.00
81 ILE O	-23.00	-22.26	79.29	15.00
82 ASP N	-21.37	-20.92	78.48	15.00
82 ASP CA	-21.88	-19.75	79.15	15.00
82 ASP CB	-20.83	-18.66	79.19	15.00
82 ASP CG	-19.68	-19.00	80.10	15.00
82 ASP OD1	-18.57	-18.52	79.85	15.00
82 ASP OD2	-19.91	-19.76	81.07	15.00
82 ASP C	-23.17	-19.21	78.57	15.00
82 ASP O	-23.56	-19.54	77.45	15.00
83 SER N	-23.85	-18.42	79.39	15.00
83 SER CA	-25.09	-17.77	79.00	15.00
83 SER CB	-25.89	-17.34	80.22	15.00
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83	SER	OG	-25.29	-16.22	80.86	15.00
83	SER	С	-24.70	-16.56	78.16	15.00
83	SER	0	-23.58	-16.04	78.30	15.00
84	GLU	N	-25.59	-16.10	77.30	15.00
84	GLU	CA	-25.30	-14.93	76.49	15.00
84	GLU	CB	-26.55	-14.46	75.73	15.00
84	GLU	CG	-26.40	-13.12	75.04	15.00
84	GLU	CD	-25.37	-13.12	73.92	15.00
84	GLU	OE1	-24.73	-12.07	73.72	15.00
84	GLU	OE2	-25.20	-14.15	73.23	15.00
84	GLU	С	-24.74	-13.81	77.37	15.00
84	GLU	0	-23.72	-13.20	77.06	15.00
85	ASP	N	-25.37	-13.60	78.53	15.00
85	ASP	CA	-24.94	-12.54	79.44	15.00
85	ASP	CB	-25.85	-12.44	80.66	15.00
85	ASP	CG	-27.22	-11.88	80.32	15.00
85	ASP	OD1	-28.19	-12.27	81.01	15.00
85	ASP	OD2	-27.33	-11.06	79.37	15.00
85	ASP	C	-23.49	-12.60	79.87	15.00
85	ASP	0	-22.78	-11.59	79.79	15.00
86	ALA	N	-23.05	-13.79	80.28	15.00
86	ALA	CA	-21.67	-14.00	80.73	15.00
86	ALA	CB	-21.59	-15.24	81.59	15.00
86	ALA	С	-20.61	-14.06	79.62	15.00
86	ALA	0	-19.42	-14.18	79.92	15.00
87	TYR	N	-21.04	-13.98	78.36	15.00
87	TYR	CA	-20.13	-14.04	77.22	15.00
87	TYR	СВ	-19.69	-15.50	76.99	15.00
87	TYR		-18.30	-15.71	76.41	15.00
87	TYR		-17.79	-14.89	75.41	15.00
	TYR		-16.54	-15.12	74.86	15.00
87	TYR		-17.51	-16.78	76.84	15.00
87	TYR		-16.26	-17.02	76.29	15.00
87	TYR		-15.78	-16.18	75.30	15.00
87	TYR		-14.54	-16.42	74.76	15.00
87	TYR		-20.88	-13.46	76.00	15.00
87	TYR		-21.14	-14.15	75.02	15.00
88	PRO		-21.17	-12.14	76.05	15.00
88	PRO		-20.72	-11.15	77.06	15.00
88	PRO		-21.90	-11.45	74.97	15.00
88	PRO		-21.94	-9.99	75.46	15.00
88	PRO	CG	-20.72	-9.87	76.27	15.00

88	PRO C	-21.30	-11.57	73.58	15.00
88	PRO C	-20.11	-11.80	73.42	15.00
89	TYR N	-22.15	-11.39	72.58	15.00
89	TYR CA	-21.74	-11.48	71.18	15.00
89	TYR CB	-22.93	-11.78	70.27	15.00
89	TYR CG	-22.53	-12.08	68.84	15.00
89	TYR CD1	-21.82	-13.23	68.54	15.00
89	TYR CE1	-21.40	-13.50	67.25	15.00
89	TYR CD2	-22.83	-11.20	67.81	15.00
89	TYR CE2	-22.40	-11.46	66.50	15.00
89	TYR CZ	-21.69	-12.62	66.24	15.00
89	TYR OH	-21.23	-12.90	64.98	15.00
89	TYR C	-21.04	-10.21	70.71	15.00
89	TYR O	-21.54	-9.11	70.94	15.00
90	VAL N	-19.88	-10.36	70.08	15.00
90	VAL CA	-19.15	-9.21	69.55	15.00
90	VAL CB	-17.81	-8.94	70.27	15.00
90	VAL CG1	-18.06	-8.15	71.55	15.00
90	VAL CG2	-17.08	-10.23	70.56	15.00
90	VAL C	-18.92	-9.31	68.05	15.00
90	VAL O	-18.60	-8.32	67.40	15.00
91	GLY N	-19.08	-10.51	67.50	15.00
91	GLY CA	-18.90	-10.68	66.06	15.00
91	GLY C	-17.46	-10.67	65.56	15.00
91	GLY O	-17.19	-10.28	64.42	15.00
92	GLN N	-16.54	-11.10	66.41	15.00
92	GLN CA	-15.14	-11.17	66.04	15.00
92	GLN CB	-14.46	-9.80	66.11	15.00
92	GLN CG	-14.41	-9.16	67.49	15.00
92	GLN CD	-14.16	-7.65	67.45	15.00
92	GLN OE1	-14.60	-6.91	68.33	15.00
92	GLN NE2	-13.46	-7.18	66.42	15.00
92	GLN C	-14.45	-12.22	66.92	15.00
92	GLN O	-14.82	-12.42	68.07	15.00
93	GLU N	-13.51	-12.94	66.32	15.00
93	GLU CA	-12.75	-13.98	66.99	15.00
93	GLU CB	-11.92	-14.80	65.98	15.00
93	GLU CG	-12.60	-15.10	64.64	15.00
93	GLU CD	-11.66	-15.74	63.60	15.00
93	GLU OE1	-10.42	-15.60	63.72	15.00
93	GLU OE2	-12.17	-16.37	62.65	15.00
93	GLU C	-11.83	-13.34	68.03	15.00

93	GLU	0	-11.21	-12.30	67.75	15.00
94	GLU	N	-11.73	-13.97	69.20	15.00
94	GLU	CA	-10.88	-13.49	70.30	15.00
94	GLU	CB	-11.58	-12.42	71.15	15.00
94	GLU	CG	-11.71	-11.07	70.44	15.00
94	GLU	CD	-12.53	-10.04	71.21	15.00
94	GLU	OE1	-13.43	-10.42	71.98	15.00
94	GLU	OE2	-12.26	-8.84	71.02	15.00
94	GLU	С	-10.47	-14.70	71.15	15.00
94	GLU	0	-10.92	-15.82	70.89	15.00
95	SER	N	-9.61	-14.49	72.13	15.00
95	SER	CA	-9.17	-15.58	72.99	15.00
95	SER	CB	-7.98	-15.16	73.87	15.00
95	SER	OG	-8.29	-14.05	74.70	15.00
95	SER	C	-10.29	-16.20	73.83	15.00
95	SER	0	-11.16	-15.50	74.36	15.00
96	CYS	N	-10.27	-17.53	73.93	15.00
96	CYS	CA	-11.26	-18.23	74.72	15.00
96	CYS	C	-11.28	-17.62	76.13	15.00
96	CYS	0	-10.26	-17.58	76.83	15.00
96	CYS	СВ	-10.97	-19.73	74.75	15.00
96	CYS	SG	-12.23	-20.69	75.64	15.00
97	MET	N	-12.44	-17.08	76.50	15.00
97	MET	CA	-12.64	-16.44	77.80	15.00
97	MET	CB	-12.80	-14.93	77.61	15.00
97	MET	CG	-12.60	-14.11	78.87	15.00
97	MET	SD	-10.92	-14.28	79.50	15.00
97	MET	CE	-10.06	-13.18	78.42	15.00
97	MET	С	-13.84	-17.05	78.53	15.00
97	MET	0	-14.64	-16.34	79.14	15.00
98	TYR		-13.96	-18.37	78.44	15.00
	TYR		-15.04	-19.09	79.10	15.00
	TYR		-15.03	-20.57	78.73	15.00
98	TYR	CG	-15.99	-21.40	79.55	15.00
	TYR		-17.36	-21.36	79.31	15.00
98	TYR		-18.25	-22.09	80.09	15.00
98	TYR		-15.53	-22.21	80.59	15.00
	TYR		-16.41	-22.95	81.38	15.00
98	TYR		-17.77	-22.88	81.13	15.00
98	TYR		-18.64	-23.57	81.93	15.00
98	TYR		-15.01	-18.91	80.61	15.00
98	TYR	0	-14.00	-19.18	81.26	15.00

99	ASN		-16.12	-18.44	81.16	15.00
99	ASN	I CA	-16.24	-18.23	82.58	15.00
99	ASN	CB	-16.80	-16.84	82.89	15.00
99	ASN	CG	-16.73	-16.50	84.38	15.00
99	ASN	OD1	-16.91	-17.36	85.24	15.00
99	ASN	ND2	-16.44	-15.25	84.68	15.00
99	asn	C	-17.14	-19.31	83.15	15.00
99	asn	0	-18.33	-19.37	82.85	15.00
100	PRO	N	-16.59	-20.19	83.99	15.00
100	PRO	CD	-15.16	-20.25	84.37	15.00
100	PRO	CA	-17.34	-21.29	84.62	15.00
100	PRO	CB	-16.27	-21.98	85.47	15.00
100	PRO	CG	-15.00	-21.70	84.70	15.00
100	PRO	С	-18.52	-20.82	85.48	15.00
100	PRO	0	-19.53	-21.51	85.58	15.00
101	THR	N	-18.37	-19.64	86.09	15.00
101	THR	CA	-19.42	-19.05	86.93	15.00
101	THR	CB	-18.92	-17.73	87.61	15.00
101	THR	OG1	-17.73	-17.97	88.38	15.00
101	THR	CG2	-19.99	-17.15	88.54	15.00
101	THR	С	-20.68	-18.73	86.12	15.00
101	THR	0	-21.77	-18.69	86.68	15.00
102	GLY	N	-20.52	-18.51	84.81	15.00
102	GLY	CA	-21.67	-18.18	83.97	15.00
102	GLY	C	-22.36	-19.33	83.25	15.00
102	GLY	0	-23.34	-19.12	82.53	15.00
103	LYS	N	-21.87	-20.54	83.47	15.00
103	LYS	CA	-22.41	-21.74	82.83	15.00
103	LYS	CB	-21.73	-22.98	83.40	15.00
103	LYS		-21.93	-24.24	82.59	15.00
103	LYS	CD	-21.93	-25.43	83.52	15.00
	LYS		-20.80	-25.36	84.52	15.00
	LYS		-21.18	-26.01	85.80	15.00
103	LYS	С	-23.91	-21.86	82.95	15.00
103			-24.44	-21.97	84.05	15.00
	ALA		-24.60	-21.92	81.82	15.00
104	ALA	CA	-26.05	-22.02	81.81	15.00
104	ALA	CB	-26.65	-20.90	80.97	15.00
104	ALA	C.	-26.59	-23.38	81.35	15.00
104	ALA	0	-27.77	-23.67	81.53	15.00
105	ALA	N	-25.72	-24.20	80.77	15.00
105	ALA	CA	-26.11	-25.53	80.29	15.00

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105	ALA	CB	-27.03	-25.41	79.07	15.00
105	ALA	С	-24.88	-26.34	79.91	15.00
105	ALA	0	-23.75	-25.83	79.94	15.00
106	LYS	N	-25.12	-27.60	79.57	15.00
106	LYS	CA	-24.10	-28.54	79.11	15.00
106	LYS	CB	-23.33	-29.21	80.26	15.00
106	LYS	CG	-24.14	-30.16	81.14	15.00
106	LYS	CD	-23.57	-31.59	81.11	15.00
106	LYS	CE	-24.32	-32.54	82.06	15.00
106	LYS	NZ	-25.78	-32.68	81.70	15.00
106	LYS	C	-24.83	-29.57	78.27	15.00
106	LYS	0	-26.05	-29.54	78.19	15.00
107	CYS	N	-24.09	-30.44	77.60	15.00
107	CYS	CA	-24.70	-31.48	76.79	15.00
	CYS		-24.95	-31.05	75.34	15.00
	CYS		-23.54	-31.08	74.19	15.00
107	CYS	С	-23.84	-32.72	76.88	15.00
107	CYS	0	-22.66	-32.64	77.19	15.00
	ARG		-24.45	-33.87	76.68	15.00
	ARG		-23.73	-35.13	76.74	15.00
108	ARG	CB	-24.36	-36.04	77.78	15.00
	ARG		-24.14	-35.57	79.21	15.00
	ARG		-24.54	-36.64	80.19	15.00
	ARG		-25.96	-36.93	80.10	15.00
	ARG		-26.54	-38.01	80.61	15.00
	ARG		-25.81	-38.91	81.24	15.00
	ARG		-27.85	-38.18	80.50	15.00
	ARG		-23.58	-35.82	75.39	15.00
	ARG		-23.92	~36.99	75.24	15.00
	GLY		-23.09	-35.08	74.41	15.00
	GLY		-22.90	-35.66	73.10	15.00
	GLY		-23.96	-35.21	72.13	15.00
	GLY			-34.38	72.48	15.00
110			-23.95	-35.79	70.94	15.00
110			-24.90	-35.45	69.88	15.00
110	TYR		-24.35	-34.33	68.98	15.00
	TYR			-34.72	68.26	15.00
	TYR			-34.52	68.86	15.00
	TYR			-34.96	68.26	15.00
110	TYR		-23.13	-35.37	67.04	15.00
110	TYR		-21.98	-35.82	66.43	15.00
110	TYR	CZ	-20.76	-35.62	67.05	15.00

110 TYR	HO	-19.61	-36.09	66.45	15.00
110 TYR	C	-25.28	-36.66	69.02	15.00
110 TYR	. 0	-24.66	-37.72	69.09	15.00
111 ARG	N	-26.28	-36.45	68.18	15.00
111 ARG	CA	-26.77	-37.47	67.27	15.00
111 ARG	CB	-28.06	-38.10	67.78	15.00
111 ARG	CG	-29.00	-38.71	66.72	15.00
111 ARG	CD	-28.59	-40.10	66.19	15.00
111 ARG	NE	-29.56	-40.59	65.20	15.00
111 ARG	CZ	-29.24	-41.12	64.02	15.00
111 ARG	NH1	-27.97	-41.25	63.65	15.00
111 ARG	NH2	-30.20	-41.45	63.15	15.00
111 ARG	C	-26.95	-36.78	65.92	15.00
111 ARG	0	-27.32	-35.60	65.85	15.00
112 GLU	N	-26.60	-37.49	64.86	15.00
112 GLU	CA	-26.73	-36.98	63.50	15.00
112 GLU	CB	-25.44	-37.25	62.71	15.00
112 GLU	CG	-24.23	-36.54	63.29	15.00
112 GLU	CD	-22.94	-36.81	62.52	15.00
112 GLU	OE1	-22.55	-36.00	61.66	15.00
112 GLU	OE2	-22.30	-37.84	62.82	15.00
112 GLU	C	-27.95	-37.64	62.84	15.00
112 GLU	0	-28.32	-38.74	63.20	15.00
113 ILE	N	-28.60	-36.94	61.93	15.00
113 ILE		-29.75	-37.51	61.24	15.00
113 ILE		-30.79	-36.40	60.90	15.00
113 ILE	CG2	-31.82	-36.90	59.89	15.00
113 ILE	CG1	-31.47	-35.89	62.17	15.00
113 ILE	CD1	-32.11	-36.98	63.00	15.00
113 ILE	С	-29.23	-38.17	59.97	15.00
113 ILE	0	-28.24	-37.70	59.39	15.00
114 PRO		-29.81	-39.32	59.56	15.00
114 PRO		-30.91	-40.08	60.16	15.00
114 PRO		-29.34	-39.97	58.34	15.00
114 PRO		-30.43	-41.01	58.08	15.00
114 PRO	CG	-30.84	-41.40	59.42	15.00
114 PRO	С	-29.30	-38.94	57.22	15.00
114 PRO	_	-30.29	-38.23	56.97	15.00
115 GLU		-28.14	-38.81	56.59	15.00
115 GLU		-27.95	-37.85	55.52	15.00
115 GLU		-26.52	-37.88	55.00	15.00
115 GLU	CG	-26.24	-36.84	53.95	15.00

115 GLU CD	-24.87	-36.97	53.34	15.00
115 GLU OE1	-24.73	-37.74	52.35	
115 GLU OE2	-23.94		53.84	
115 GLU C	-28.94	-38.05	54.38	15.00
115 GLU O	-29.14	-39.17	53.91	15.00
116 GLY N	-29.55	-36.96	53.95	15.00
116 GLY CA	-30.51	-37.02	52.86	
116 GLY C	-31.93	-37.43	53.23	15.00
116 GLY O	-32.85	-37.23	52.44	15.00
117 ASN N	-32.12	-37.92	54.46	15.00
117 ASN CA	-33.43	-38.37	54.93	15.00
117 ASN CB	-33.27	-39.59	55.85	15.00
117 ASN CG	-34.53	-40.48	55.92	15.00
117 ASN OD1	-35.65	-40.02	55.70	15.00
117 ASN ND2	-34.34	-41.74	56.27	15.00
117 ASN C	-34.30	-37.30	55.60	15.00
117 ASN 0	-34.12	-37.00	56.79	
118 GLU N	-35.26	-36.75	54.86	15.00
118 GLU CA	-36.18	-35.73	55.38	15.00
118 GLU CB	-36.91	-34.98	54.26	15.00
118 GLU CG	-36.01	-34.11	53.37	15.00
118 GLU CD	-36.78		52.55	15.00
118 GLU OE1	-36.86	-33.26	51.32	15.00
118 GLU OE2	-37.30	-32.11	53.13	15.00
118 GLU C	-37.19	-36.37	56.35	15.00
118 GLU O	-37.57	-35.76	57.35	15.00
119 LYS N	-37.59	-37.60	56.06	15.00
119 LYS CA	-38.53		56.92	15.00
119 LYS CB	-38.89	-39.68	56.33	15.00
119 LYS CG	-40.10	-40.36	56.97	15.00
119 LYS CD	-40.37	-41.75	56.35	15.00
119 LYS CE	39.71		57.16	15.00
119 LYS NZ	-39.80		56.47	15.00
119 LYS C	-37.89	-38.45	58.30	15.00
119 LYS O	-38.53	-38.23	59.33	15.00
120 ALA N	-36.59	-38.76	58.30	15.00
120 ALA CA	-35.85	-38.91	59.54	15.00
120 ALA CB	-34.44	-39.41	59.25	15.00
120 ALA C	-35.78	-37.58	60.26	15.00
120 ALA O	-35.89	-37.52	61.49	15.00
121 LEU N	-35.61	-36.51	59.48	15.00
121 LEU CA	-35.52	-35.16	60.02	15.00

	LEU		-35.02		58.94	15.00
121		CG	-34.84	-32.68	59.23	15.00
121	LEU	CD1	-33.98	-32.43	60.46	15.00
121	LEU	CD2	-34.25	-32.02	57.99	15.00
121	LEU	С	-36.83	-34.68	60.68	15.00
121	LEU	0	-36.79	-34.05	61.74	15.00
122	LYS	N	-37.97	-35.03	60.08	15.00
122	LYS	CA	-39.29	-34.65	60.60	15.00
122	LYS	CB	-40.42	-34.99	59.63	15.00
122	LYS	CG	-41.82	-34.90	60.26	15.00
122	LYS	CD	-42.89	-35.49	59.34	15.00
122	LYS	CE	-44.28	-35.44	59.97	15.00
122	LYS	NZ	-45.36	-35.91	59.03	15.00
122	LYS	С	-39.57	-35.33	61.92	15.00
122	LYS	0	-40.03	-34.70	62.88	15.00
123	ARG	N	-39.33	-36.63	61.96	15.00
	ARG		-39.55	-37.40	63.17	15.00
	ARG	CB	-39.26	-38.88	62.91	15.00
	ARG		-40.26	-39.53	61.95	15.00
	ARG		-40.04	-41.04	61.79	15.00
	ARG		-38.71	-41.35	61.25	15.00
123			-37.76	-42.03	61.89	15.00
123	ARG		-37.99	-42.51	63.12	15.00
123	ARG		-36.57	-42.20	61.33	15.00
	ARG		-38.72	-36.82	64.33	15.00
	ARG		-39.21	-36.72	65.46	15.00
	ALA		-37.50	-36.38	64.04	15.00
	ALA		-36.62	-35.79	65.05	15.00
	ALA		-35.21	-35.60	64.50	15.00
	ALA		-37.17	-34.46	65.58	15.00
	ALA		-37.17	-34.21	66.79	15.00
	VAL		-37.64	-33.61	64.68	15.00
	VAL		-38.20	-32.33	65.09	15.00
	VAL		-38.57			15.00
	VAL		-39.39		64.31	15.00
			-37.30		63.15	15.00
	VAL		-39.41		65.99	15.00
125	VAL			-31.99	67.07	15.00
126	ALA			-33.49	65.58	15.00
	ALA			-33.83	66.34	15.00
	ALA		-42.27		65.62	15.00
126	ALA	С	-41.15	-34.29	67.77	15.00

126	ALA	0	-41.69	-33.74	68.73	15.00
127	ARG	N	-40.26	-35.26	67.93	15.00
127	ARG	CA	-39.95	-35.76	69.27	15.00
127	ARG	CB	-39.99	-37.28	69.30	15.00
127	ARG	CG	-38.95	-37.98	68.45	15.00
127	ARG	CD	-39.08	-39.48	68.63	15.00
127	ARG	NE	-40.43	-39.95	68.30	15.00
127	ARG	CZ	-41.25	-40.58	69.14	15.00
127	ARG	NH1	-42.47	-40.94	68.73	15.00
127	ARG	NH2	-40.88	-40.83	70.39	15.00
127	ARG	С	-38.72	-35.26	70.01	15.00
127	ARG	0	-38.50	-35.65	71.15	15.00
128	VAL	N	-37.91	-34.42	69.40	15.00
128	VAL	CA	-36.72	-33.94	70.07	15.00
128	VAL	CB	-35.46	-34.31	69.27	15.00
128	VAL	CG1	-34.25	-33.52	69.74	15.00
128	VAL	CG2	-35.18	-35.79	69.42	15.00
128	VAL	С	-36.78	-32.43	70.32	15.00
128	VAL	0	-36.54	-31.96	71.43	15.00
129	GLY	N	-37.12	-31.68	69.28	15.00
129	GLY	CA	-37.18	-30.24	69.39	15.00
129	GLY	С	-36.41	-29.69	68.22	15.00
129	GLY	0	-36.10	-30.46	67.30	15.00
130	PRO	N	-36.09	-28.39	68.19	15.00
130	PRO	CD	-36.46	-27.40	69.22	15.00
130	PRO	CA	-35.34	-27.75	67.11	15.00
130	PRO	CB	-35.01	-26.38	67.70	15.00
130	PRO	CG	-36.23	-26.08	68.51	15.00
130	PRO	С	-34.06	-28.52	66.73	15.00
130	PRO	0	-33.35	-29.05	67.61	15.00
131	VAL	N	-33.78	-28.58	65.44	15.00
131	VAL	CA	-32.61	-29.30	64.92	15.00
131	VAL	СВ	-33.05	-30.57	64.15	15.00
131	VAL	CG1	-31.85	-31.31	63.57	15.00
131	VAL	CG2	-33.84	-31.50	65.07	15.00
131	VAL	С	-31.71	-28.42	64.02	15.00
131	VAL	0	-32.21	-27.71	63.14	15.00
132	SER	N	-30.41	-28.44	64.29	15.00
132	SER	CA	-29.40	-27.69	63.53	15.00
132	SER	СВ	-28.07	-27.65	64.28	15.00
132	SER	OG	-28.22	-27.17	65.59	15.00
132	SER	С	-29.19	-28.31	62.13	15.00

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132 SER O	-28.82	-29.48	62.02	15.00
133 VAL N	-29.40	-27.52	61.08	15.00
133 VAL CA	-29.23	-28.02	59.73	15.00
133 VAL CB	-30.60	-28.18	58.99	15.00
133 VAL CG1	-31.53	-29.06	59.80	15.00
133 VAL CG2	-31.24	-26.84	58.70	15.00
133 VAL C	-28.35	-27.10	58.89	15.00
133 VAL 0	-28.22	-25.92	59.20	15.00
134 ALA N	-27.74	-27.66	57.85	15.00
134 ALA CA	-26.88	-26.90	56.95	15.00
134 ALA CB	-25.50	-27.56	56.83	15.00
134 ALA C	-27.59	-26.86	55.59	15.00
134 ALA O	-28.15	-27.87	55.15	15.00
135 ILE N	-27.61	-25.69	54.96	15.00
135 ILE CA	-28.28	-25.52	53.68	15.00
135 ILE CB	-29.64	-24.75	53.86	15.00
135 ILE CG2	-30.59	-25.51	54.77	15.00
135 ILE CG1	-29.37	-23.34	54.39	15.00
135 ILE CD1	-30.61	-22.47	54.50	15.00
135 ILE C	-27.45	-24.69	52.71	15.00
135 ILE 0	-26.36	-24.22	53.04	15.00
136 ASP N	-27.98	-24.56	51.49	15.00
136 ASP CA	-27.37	-23.75	50.45	15.00
136 ASP CB	-27.45	-24.42	49.07	15.00
136 ASP CG	-26.86	-23.57	47.94	15.00
136 ASP OD1	-26.91	-24.02	46.79	15.00
136 ASP OD2	-26.35	-22.45	48.19	15.00
136 ASP C	-28.21	-22.46	50.50	15.00
136 ASP 0	-29.41	-22.48	50.22	15.00
137 ALA N	-27.58	-21.38	50.92	15.00
137 ALA CA	-28.23	-20.08	51.04	15.00
137 ALA CB	-28.30	-19.68	52.49	15.00
137 ALA C	-27.45	-19.04	50.25	15.00
137 ALA O	-27.31	-17.91	50.69	15.00
138 SER N	-26.97	-19.44	49.08	15.00
138 SER CA	-26.18	-18.56	48.22	15.00
138 SER CB	-25.05	-19.36	47.56	15.00
138 SER OG	-25.57	-20.33	46.67	15.00
138 SER C	-26.99	-17.81	47.16	15.00
138 SER O		-16.88	46.52	15.00
139 LEU N	-28.23	-18.24	46.97	15.00
139 LEU CA	-29.11	-17.68	45.97	15.00

139	LEU	CB	-30.17	-18.70	45.55	15.00
139	LEU	CG	-29.68	-19.89	44.71	15.00
139	LEU	CD1	-28.41	-20.51	45.27	15.00
139	LEU	CD2	-30.78	-20.93	44.65	15.00
139	LEU	С	-29.76	-16.35	46.31	15.00
139	LEU	0	-30.20	-16.13	47.44	15.00
140	THR	N	-29.82	-15.48	45.31	15.00
140	THR	CA	-30.41	-14.15	45.44	15.00
140	THR	CB	-30.31	-13.40	44.07	15.00
140	THR	OG1	-28.94	-13.06	43.81	15.00
140	THR	CG2	-31.17	-12.14	44.06	15.00
140	THR	С	-31.87	-14.27	45.91	15.00
140	THR	0	-32.36	-13.41	46.66	15.00
141	SER	N	-32.55	-15.34	45.50	15.00
141	SER	CA	-33.93	-15.58	45.89	15.00
141	SER	CB	-34.47	-16.84	45.23	15.00
141	SER	OG	-33.61	-17.94	45.44	15.00
141	SER	С	-34.00	-15.70	47.41	15.00
141	SER	0	-34.83	-15.07	48.07	15.00
142	PHE	N	-33.07	-16.47	47.97	15.00
142	PHE	CA	-32.99	-16.70	49.40	15.00
142	PHE	CB	-31.94	-17.76	49.73	15.00
142	PHE	CG	-31.90	-18.15	51.17	15.00
142	PHE	CD1	-32.77	-19.12	51.66	15.00
142	PHE	CD2	-31.02	-17.55	52.05	15.00
142	PHE	CE1	-32.75	-19.48	53.00	15.00
142	PHE	CE2	-31.00	-17.90	53.39	15.00
142	PHE	CZ	-31.86	-18.87	53.86	15.00
142	PHE	С	-32.73	-15.42	50.16	15.00
142	PHE	0	-33.46	-15.07	51.08	15.00
143	GLN	N	-31.67	-14.71	49.76	15.00
143	GLN	CA	-31.30	-13.45	50.40	15.00
	GLN		-30.10	-12.81	49.70	15.00
143	GLN	CG	-29.69	-11.45	50.28	15.00
143	GLN	CD	-28.34	-10.91	49.77	15.00
143	GLN	OE1	-27.73	-10.04	50.40	15.00
143	GLN	NE2	-27.85	-11.46	48.64	15.00
	GLN	С	-32.50	-12.51	50.43	15.00
143	GLN	0	-32.76	-11.85	51.44	15.00
144	PHE	N	-33.28	-12.51	49.36	15.00
144	PHE	CA	-34.43	-11.62	49.26	15.00
144	PHE	CB	-34.43	-10.90	47.89	15.00

144 PHE CG	-33.21	-10.01	47.66	15.00
144 PHE CD1	-32.96	-8.92	48.48	15.00
144 PHE CD2	-32.31	-10.29	46.63	15.00
144 PHE CE1	-31.83	-8.12	48.30	15.00
144 PHE CE2	-31.17	-9.48	46.44	15.00
144 PHE CZ	-30.94	-8.40	47.27	15.00
144 PHE C	-35.79	-12.23	49.61	15.00
144 PHE O	-36.83	-11.62	49.35	15.00
145 TYR N	-35.78	-13.41	50.24	15.00
145 TYR CA	-37.03	-14.08	50.64	15.00
145 TYR CB	-36.76	-15.42	51.35	15.00
145 TYR CG	-37.96	-16.00	52.08	15.00
145 TYR CD1	-38.82	-16.92	51.46	15.00
145 TYR CE1	-39.93	-17.45	52.13	15.00
145 TYR CD2	-38.26	-15.62	53.38	15.00
145 TYR CE2	-39.37	-16.13	54.05	15.00
145 TYR CZ	-40.20	-17.04	53.43	15.00
145 TYR OH	-41.29	-17.54	54.12	15.00
145 TYR C	-37.90	-13.18	51.52	15.00
145 TYR O	-37.38	-12.43	52.36	15.00
146 SER N	-39.21	-13.31	51.38	15.00
146 SER CA	-40.13	-12.49	52.15	15.00
146 SER CB	-40.37	-11.15	51.45	15.00
146 SER OG	-40.91	-11.35	50.15	15.00
146 SER C	-41.46	-13.15	52.51	15.00
146 SER O	-41.95	-12.96	53.62	15.00
147 LYS N	-42.01	-13.92	51.58	15.00
147 LYS CA	-43.30	-14.59	51.76	15.00
147 LYS CB	-44.42	-13.76	51.10	15.00
147 LYS CG	-44.60	-12.34	51.62	15.00
147 LYS CD	-45.05	-12.34	53.08	15.00
147 LYS CE	-45.37	-10.92	53.58	15.00
147 LYS NZ	-46.00	-10.90	54.93	15.00
147 LYS C		-16.03	51.20	15.00
147 LYS O	-42.63	-16.39	50.27	15.00
148 GLY N	-44.29	-16.81	51.75	15.00
148 GLY CA	-44.52	-18.16	51.28	15.00
148 GLY C	-43.46	-19.22	51.49	15.00
148 GLY 0	-42.47	-18.99	52.18	15.00
149 VAL N	-43.70	-20.41	50.95	15.00
149 VAL CA		-21.53	51.07	15.00
149 VAL CB	-43.53	-22.90	50.95	15.00

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149	VAL	CG1	-42.55	-24.07	50.90	15.00
149	VAL	CG2	-44.47	-23.08	52.14	15.00
149	VAL	С	-41.69	-21.41	50.01	15.00
149	VAL	0	-41.94	-21.55	48.82	15.00
150	TYR	N	-40.49	-21.08	50.48	15.00
150	TYR	CA	-39.31	-20.92	49.63	15.00
150	TYR	CB	-38.12	-20.36	50.42	15.00
150	TYR	CG	-36.84	-20.29	49.60	15.00
150	TYR	CD1	-36.67	-19.30	48.63	15.00
150	TYR	CE1	-35.54	-19.29	47.81	15.00
150	TYR	CD2	-35.84	-21.25	49.74	15.00
150	TYR	CE2	-34.71	-21.24	48.92	15.00
150	TYR	CZ	-34.57	-20.26	47.96	15.00
150	TYR	OH	-33.48	-20.27	47.12	15.00
150	TYR	C	-38.89	-22.18	48.89	15.00
150	TYR	0	-38.88	-23.28	49.45	15.00
151	TYR		-38.47	-21.98	47.65	15.00
151	TYR	CA	-37.98	-23.03	46.77	15.00
151	TYR	CB	-39.09	-23.99	46.35	15.00
	TYR		-38.62	-25.09	45.42	15.00
	TYR		-37.92	-26.20	45.91	15.00
	TYR	CE1	-37.51	-27.23	45.06	15.00
151		CD2	-38.89	-25.04	44.05	15.00
151	TYR	CE2	-38.49	-26.07	43.19	15.00
151	TYR	CZ	-37.80	-27.16	43.70	15.00
151	TYR	OH	-37.46	-28.21	42.87	15.00
151	TYR	С	-37.35	-22.39	45.55	15.00
151	TYR		-37.80	-21.33	45.07	15.00
	ASP		-36.30	-23.02	45.05	15.00
	ASP		-35.59	-22.54	43.86	15.00
	ASP		-34.66	-21.38	44.20	15.00
	ASP		-34.13	-20.68	42.97	15.00
	ASP		-33.52	-19.60	43.12	15.00
152	ASP	OD2	-34.32	-21.20	41.84	15.00
	ASP			-23.70	43.25	15.00
	ASP			-24.27	43.89	15.00
	GLU			-24.04	42.01	15.00
	GLU			-25.15	41.34	15.00
153				-25.45	40.00	15.00
153	GLU			-24.38	38.96	15.00
	GLU			-24.81	37.56	15.00
153	GLU	OE1	-35.72	-26.00	37.37	15.00

153 GLU OE2	35 40			
153 GLU C		-23.94		
153 GLU 0	-32.98		41.17	15.00
	-32.30		40.92	15.00
	-32.46	-23.80	41.32	15.00
154 SER CA	-31.02			15.00
154 SER CB	-30.77		40.66	15.00
154 SER OG	-31.56		39.50	15.00
154 SER C	-30.23		42.46	15.00
154 SER O	-28.99		42.45	15.00
155 CYS N	-30.94		43.53	15.00
155 CYS CA	-30.35		44.84	15.00
155 CYS C	-29.60	-25.71	44.92	15.00
155 CYS 0	-30.20	-26.78	44.85	15.00
155 CYS CB	-31.43	-24.32	45.91	15.00
155 CYS SG	-30.84	-23.76	47.53	15.00
156 ASN N	-28.29	-25.64	45.11	15.00
156 ASN CA	-27.46	-26.84	45.20	15.00
156 ASN CB	-26.08	-26.58	44.61	15.00
156 ASN CG	-25.26	-27.85	44.48	15.00
156 ASN OD1	-25.77	-28.97	44.56	15.00
156 ASN ND2	-23.96	-27.69	44.26	15.00
156 ASN C	-27.33	-27.51	46.58	15.00
156 ASN 0	-26.74	-26.95	47.51	15.00
157 SER N	-27.78	-28.76	46.65	15.00
157 SER CA	-27.73	-29.53	47.88	15.00
157 SER CB	-28.66	-30.74	47.80	15.00
157 SER OG	-28.22	-31.64	46.80	15.00
157 SER C	-26.33	-29.97	48.29	15.00
157 SER O	-26.16	-30.58	49.35	15.00
158 ASP N	-25.34	-29.71	47.43	15.00
158 ASP CA	-23.95	-30.07	47.71	15.00
158 ASP CB	-23.35	-30.92	46.59	15.00
158 ASP CG	-24.02	-32.27	46.47	15.00
158 ASP OD1	-24.01	-33.04	47.47	15.00
158 ASP OD2	-24.58	-32.55	45.38	15.00
158 ASP C	-23.09	-28.84	47.97	15.00
158 ASP 0	-21.86	-28.90	47.96	15.00
159 ASN N	-23.76	-27.70	48.14	15.00
159 ASN CA			48.42	15.00
159 ASN CB		-25.45	47.30	15.00
159 ASN CG		-24.14	47.51	15.00
159 ASN OD1		-24.00	48.41	15.00
			<b></b>	

159	ASN	ND2	-22.86	-23.16	46.67	15.00
159	ASN	C	-23.69	-25.99	49.73	15.00
159	ASN	0	-24.50	-25.06	49.75	15.00
160	LEU	N	-23.39	-26.71	50.81	15.00
160	LEU	CA	-23.89	-26.38	52.15	15.00
160	LEU	CB	-23.90	-27.62	53.06	15.00
160	LEU	CG	-24.57	-28.92	52.58	15.00
160	LEU	CD1	-24.38	-30.01	53.60	15.00
160	LEU	CD2	-26.04	-28.72	52.33	15.00
160	LEU	C	-23.01	-25.25	52.70	15.00
160	LEU	0	-21.92	-25.48	53.22	15.00
161	ASN	N	-23.52	-24.02	52.60	15.00
161	asn	CA	-22.76	-22.83	53.00	15.00
161	asn	CB	-22.57	-21.92	51.79	15.00
161	asn	CG	-23.88	-21.63	51.05	15.00
161	asn	OD1	-24.76	-20.95	51.55	15.00
161	asn	ND2	-24.00	-22.18	49.86	15.00
161	asn	С	-23.26	-22.01	54.20	15.00
161	asn	0	-22.55	-21.12	54.69	15.00
162	HIS	N	-24.46	-22.32	54.68	15.00
	HIS		-25.05	-21.57	55.79	15.00
	HIS		-25.96	-20.49	55.23	15.00
	HIS		-26.59	-19.59	56.25	15.00
	HIS		-27.87	-19.21	56.42	15.00
	HIS		-25.86	-18.91	57.21	15.00
	HIS		-26.66	-18.14	57.91	15.00
	HIS	NE2	-27.89	-18.30	57.45	15.00
162	HIS		-25.78	-22.50	56.78	15.00
	HIS		-26.69	-23.22	56.39	15.00
	ALA		-25.32	-22.55	58.02	15.00
			-25.96	-23.37	59.05	15.00
	ALA		-24.98	-23.74	60.15	15.00
	ALA			-22.57	59.62	15.00
	ALA	_	-26.96		60.03	15.00
	VAL		-28.29		59.71	15.00
	VAL			-22.54	60.16	15.00
	VAL			-22.26	58.88	15.00
	VAL		-31.36	-23.35	58.63	15.00
	VAL		-30.92	-20.89	58.89	15.00
	VAL			-23.45	61.21	15.00
	VAL		-29.65		61.57	15.00
165	LEU	N	-31.35	-23.06	61.74	15.00

165 LEU CA	-32.05	-23.90	62.72	15.00
165 LEU CB	-32.05	-23.25	64.11	15.00
165 LEU CG	-32.78	-23.96	65.26	15.00
165 LEU CD1	-32.01	-25.18	65.72	15.00
165 LEU CD2	-32.97	-22.99	66.42	15.00
165 LEU C	-33.48	-24.25	62.31	15.00
165 LEU O	-34.26	-23.36	61.97	15.00
166 ALA N	-33.83	-25.54	62.36	15.00
166 ALA CA	-35.17	-26.00	62.00	15.00
166 ALA CB	-35.08	-27.37	61.33	15.00
166 ALA C	-36.09	-26.07	63.23	15.00
166 ALA O	-36.08	-27.05	63.97	15.00
167 VAL N	-36.92	-25.04	63.39	15.00
167 VAL CA	-37.82	-24.95	64.54	15.00
167 VAL CB	-37.90	-23.48	65.05	15.00
167 VAL CG1	-36.52	-22.98	65.43	15.00
167 VAL CG2	-38.52	-22.57	64.00	15.00
167 VAL C	-39.23	-25.53	64.35	15.00
167 VAL 0	-40.12	-25.32	65.18	15.00
168 GLY N	-39.43	-26.27	63.27	15.00
168 GLY CA	-40.73	-26.86	63.04	15.00
168 GLY C	-40.94	-27.30	61.61	15.00
168 GLY 0	-40.00	-27.42	60.82	15.00
169 TYR N	-42.21	-27.55	61.29	15.00
169 TYR CA	-42.63	-28.00	59.98	15.00
169 TYR CB	-42.14	-29.44	59.67	15.00
169 TYR CG	-42.65	-30.54	60.59	15.00
169 TYR CD1	-43.92	-31.12	60.41	15.00
169 TYR CE1	-44.37	-32.15	61.25	15.00
169 TYR CD2	-41.86	-31.02	61.63	15.00
169 TYR CE2	-42.31	-32.06	62.47	15.00
169 TYR CZ	-43.57	-32.61	62.27	15.00
169 TYR OH	-44.00	-33.61	63.11	15.00
169 TYR C	-44.14	-27.91	59.91	15.00
169 TYR O	-44.83	-27.98	60.92	15.00
170 GLY N	-44.65	-27.78	58.70	15.00
170 GLY CA	-46.08	-27.68	58.51	15.00
170 GLY C	-46.38	-27.73	57.04	15.00
170 GLY O	-45.57	-28.17	56.24	15.00
171 ILE N	-47.52	-27.17	56.68	15.00
171 ILE CA	-47.97	-27.14	55.30	15.00
171 ILE CB	-48.97	-28.38	55.01	15.00

171	ILE	CG2	-49.85	-28.63	56.15	15.00
171	ILE	CG1	-49.58	-28.25	53.67	15.00
171	ILE	CD1	-50.27	-29.51	53.26	15.00
171	ILE	С	-48.71	-25.83	55.06	15.00
171	ILE	0	-49.51	-25.40	55.88	15.00
172	GLN	N	-48.35	-25.13	53.99	15.00
172	GLN	CA	-48.99	-23.87	53.65	15.00
172	GLN	CB	-48.01	-22.70	53.72	15.00
172	GLN	CG	-48.67	-21.36	53.62	15.00
172	GLN	CD	-47.71	-20.23	53.89	15.00
172	GLN	OE1	-47.35	-19.97	55.05	15.00
172	GLN	NE2	-47.27	-19.55	52.84	15.00
172	GLN	С	-49.60	-24.00	52.25	15.00
172	GLN	0	-48.88	-24.13	51.26	15.00
173	LYS	N	-50.93	-23.99	52.20	15.00
173	LYS	CA	-51.68	-24.15	50.97	15.00
173	LYS	CB	-51.66	-22.89	50.10	15.00
173	LYS	CG	-52.38	-23.03	48.72	15.00
173	LYS	CD	-53.92	-23.15	48.79	15.00
173	LYS	CE	-54.45	-24.44	49.47	15.00
173			-54.10	-25.73	48.79	15.00
173	LYS	C	-51.15	-25.34	50.20	15.00
173	LYS		-50.54	-25.19	49.14	15.00
174	GLY	N	-51.31	-26.53	50.77	15.00
	GLY		-50.84	-27.73	50.10	15.00
	GLY		-49.36	-27.74	49.76	15.00
174	GLY	0 .	-48.95	-28.42	48.82	15.00
175	ASN	N	-48.57	-26.97	50.50	15.00
	ASN		-47.12	-26.90	50.30	15.00
	asn		-46.69	-25.54	49.74	15.00
	ASN		-46.88	-25.43	48.24	15.00
	asn		-45.91		47.48	15.00
	ASN		-48.13	-25.38	47.79	15.00
175	asn	C	-46.40	-27.18	51.61	15.00
	asn		-46.33	-26.31	52.49	15.00
	LYS		-45.91	-28.41	51.76	15.00
	LYS			-28.82	52.97	15.00
	LYS			-30.32	52.95	15.00
	LYS			-31.18	52.92	15.00
	LYS		-45.83		52.63	15.00
	LYS			-33.51		15.00
176	LYS	NZ	-47.62	-33.57	54.13	15.00

176 LYS C	-43.90	-28.00	53.08	15.00
176 LYS 0	-43.34	-27.58	52.07	15.00
177 HIS N	-43.42	-27.80	54.30	15.00
177 HIS CA	-42.23	-26.99	54.52	15.00
177 HIS CB	-42.58	-25.51	54.38	15.00
177 HIS CG	-43.53	-25.04	55.43	15.00
177 HIS CD2	-43.36	-24.83	56.76	15.00
177 HIS ND1	-44.87	-24.83	55.18	15.00
177 HIS CE1	-45.48	-24.50	56.31	15.00
177 HIS NE2	-44.59	-24.50	57.28	15.00
177 HIS C	-41.62	-27.24	55.88	15.00
177 HIS O	-42.21	-27.92	56.71	15.00
178 TRP N	-40.49	-26.58	56.12	15.00
178 TRP CA	-39.74	-26.60	57.38	15.00
178 TRP CB	-38.28	-27.02	57.18	15.00
178 TRP CG	-38.08	-28.36	56.62	15.00
178 TRP CD2	-38.10	-29.59	57.34	15.00
178 TRP CE2	-37.84	-30.62	56.41	15.00
178 TRP CE3	-38.30	-29.94	58.68	15.00
178 THP CD1	-37.82	-28.67	55.32	15.00
178 THP NEI	-37.68	-30.02	55.18	15.00
178 TRP CZ2	-37.79	-31.98	56.77	15.00
178 TRP CZ3	-38.25	-31.28	59.05	15.00
178 TRP CH2	-37.99	-32.28	58.09	15.00
178 TRP C	-39.76	-25.15	57.89	15.00
178 TRP 0	-39.63	-24.21	57.10	15.00
179 ILE N	-39.96	-24.95	59.18	15.00
179 ILE CA	-39.96	-23.60	59.72	15.00
179 ILE CB	-40.92	-23.46	60.92	15.00
179 ILE CG2	-41.00	-22.01	61.38	15.00
179 ILE CG1	-42.33	-23.91	60.50	15.00
179 ILE CD1	-43.31	-23.94	61.64	15.00
179 ILE C	-38.50	-23.37	60.10	15.00
179 ILE 0		-24.03	61.00	15.00
180 ILE N	-37.84	-22.48	59.36	15.00
180 ILE CA	-36.43	-22.18	59.55	15.00
180 ILE CB	-35.67	-22.22	58.19	15.00
180 ILE CG2	-34.25	-21.74	58.35	15.00
180 ILE CG1	-35.73	-23.63	57.58	15.00
180 ILE CD1	-35.19	-24.73	58.48	15.00
190 IFE C	-36.14	-20.84	60.22	15.00
180 ILE O	-36.63	-19.80	59.79	15.00

181	LYS	N	-35.33	-20.88	61.27	15.00
181	LYS	CA	-34.93	-19.68	62.01	15.00
181	LYS	CB	-34.89	-19.94	63.51	15.00
181	LYS	CG	-34.35	-18.76	64.31	15.00
181	LYS	CD	-34.18	-19.11	65.77	15.00
181	LYS	CE	-33.68	-17.92	66.56	15.00
181	LYS	NZ	-33.51	-18.24	68.01	15.00
181	LYS	С	-33.55	-19.25	61.51	15.00
181	LYS	0	-32.61	-20.03	61.53	15.00
182	ASN	N	-33.43	-18.00	61.07	15.00
182	ASN	CA	-32.16	-17.49	60.56	15.00
182	ASN	CB	-32.35	-16.82	59.19	15.00
182	ASN	CG	-31.05	-16.69	58.41	15.00
182	ASN	OD1	-30.02	-17.26	58.78	15.00
182	ASN	ND2	-31.09	-15.95	57.31	15.00
182	ASN	С	-31.49	-16.53	61.56	15.00
182		0	-32.12	-16.09	62.52	15.00
183		N	-30.20	-16.28	61.35	15.00
183			-29.44	-15.37	62.20	15.00
	SER	CB	-28.30	-16.09	62.93	15.00
183			-27.53	-16.88	62.05	15.00
183			-28.93	-14.14	61.42	15.00
183	SER		<b>-27.77</b>	-13.73	61.54	15.00
184	TRP		-29.81	-13.60	60.57	15.00
184			-29.50	-12.41	59.78	15.00
184			-29.71	-12.65	58.29	15.00
184			-28.64	-13.47	57.67	15.00
184		CD2	-28.66	-14.06	56.37	15.00
184	TRP		-27.47	-14.81	56.23	15.00
	TRP		-29.58	-14.04	55,31	15.00
184	TRP		-27.46	-13.86	58.25	15.00
184		N.E.1	-26.76	-14.67	57.39	15.00
184		C2.2	-27.18	-15.53	55.06	15.00
	TRP			-14.76	54.16	15.00
	TRP			-15.49	54.04	15.00
184				-11.22	60.27	15.00
184			-30.59	-10.27	59.53	15.00
185	GLY		-30.78	-11.31	61.52	15.00
	GLY		-31.57	-10.25	62.10	15.00
	GLY			-10.45	61.90	15.00
	GLY		-33.50	-11.18		
186	GLU	N	-33.81	-9.76	62.75	15.00

186 GLU CA	-35.27			
186 GLU CB	-35.81	_	_	
186 GLU CG	-37.34	· — —		
186 GLU CD	-37.84			
186 GLU OE1	-37.53	02	65.44	
186 GLU OE2	-38.54		_	<del></del>
186 GLU C	-35.85		- <del>-</del>	15.00
186 GLU O	-36.99		61.50	15.00
187 ASN N	-35.05		61.12	15.00
187 ASN CA	-35.52	-8.27	60.84	15.00
187 ASN CB		-7.59	59.64	15.00
187 ASN CG	~35.00	-6.16	59.56	15.00
187 ASN CG	-36.02	-5.17	60.07	15.00
187 ASN ND2	-35.91	-4.68	61.20	15.00
187 ASN ND2	-37.05	-4.90	59.25	15.00
187 ASN 0	-35.23	-8.31	58.33	15.00
188 TRP N	-35.26	-7.72	57.27	15.00
188 TRP CA	-34.94	-9.60	58.43	15.00
188 TRP CB	-34.68	-10.39	57.25	15.00
188 TRP CG	-33.32	-11.10	57.33	15.00
188 TRP CD2	-33.12	-12.00	56.19	15.00
188 TRP CE2	-33.49	-13.38	56.12	15.00
188 TRP CE3	-33.22	-13.81	54.80	15.00
188 TRP CD1	-34.04	-14.29	57.03	15.00
188 TRP NE1	-32.64	-11.66	54.96	15.00
188 TRP CZ2	-32.71	-12.74	54.12	15.00
188 TRP CZ3	-33.48	-15.13	54.37	15.00
188 TRP CH2	-34.30	-15.60	56.60	15.00
188 TRP C	-34.02	-16.00	55.28	15.00
188 TRP O	-35.82	-11.39	57.10	15.00
189 GLY N	-36.41	-11.81	58.09	15.00
189 GLY CA	-36.11 -37.16	-11.76	55.85	15.00
189 GLY C	-37.16 -38.43	-12.72	55.57	15.00
189 GLY O		-12.37	56.32	15.00
190 ASN N	-38.82		56.40	15.00
190 ASN CA		-13.38	56.89	15.00
190 ASN CB		-13.15	57.64	15.00
190 ASN CB		-14.19	57.30	15.00
190 ASN CG		-13.73	57.68	15.00
190 ASN 0D1		-14.38	57.34	1.5.00
190 ASN ND2	-42.88		58.37	15.00
190 ASN C	-40.02		59.13	15.00
TO NEW O	-40.22	-14.15	59.80	15.00

191	LYS	N	-39.56	-11.99	59.63	15.00
191	LYS	CA	-39.23	-11.78	61.05	15.00
191	LYS	CB	-40.40	-12.11	61.99	15.00
191	LYS	CG	-41.62	-11.23	61.84	15.00
191	LYS	CD	-42.63	-11.52	62.95	15.00
191	LYS	CE	-43.99	-10.85	62.71	15.00
191	LYS	NZ	-44.94	-11.70	61.92	15.00
191	LYS	С	-37.98	-12.58	61.45	15.00
191	LYS	0	-37.79	-12.95	62.61	15.00
192	GLY	N	-37.13	-12.82	60.45	15.00
192	GLY	CA	-35.90	-13.56	60.66	15.00
192	GLY	С	-36.13	-15.04	60.40	15.00
192	GLY	0	-35.29	-15.88	60.72	15.00
193	TYR	N	-37.28	-15.34	59.80	15.00
193	TYR	CA	-37.66	-16.70	59.49	15.00
193	TYR	CB	-38.91	-17.11	60.25	15.00
193	TYR	CG	-38.66	-17.45	61.68	15.00
193	TYR	CD1	-38.59	-16.46	62.66	15.00
193	TYR	CE1	-38.33	-16.78	63.98	15.00
193	TYR	CD2	-38.48	-18.78	62.07	15.00
193	TYR	CE2	-38.22	-19.11	63.38	15.00
193	TYR	CZ	-38.15	-18.11	64.34	15.00
193	TYR	OH	-37.89	-18.43	65.64	15.00
193	TYR	С	-37.86	-16.93	58.01	15.00
193	TYR	0	-37.92	-15.99	57.22	15.00
194	ILE	N	-37.97	-18.20	57.65	15.00
194	ILE	CA	-38.20	-18.60	56.28	15.00
194	ILE	CB	-36.88	-18.58	55.44	15.00
194	ILE	CG2	-35.81	-19.44	56.08	15.00
194	ILE	CG1	-37.16	-19.03	54.00	15.00
194	ILE	CD1	-36.05	-18.68	53.02	15.00
194	ILE	С	-38.85	-20.00	56.28	15.00
194	ILE	0	-38.54	-20.83	57.14	15.00
195	LEU	N-	-39.84	-20.20	55.42	15.00
195	LEU	CA	-40.50	-21.49	55.30	15.00
195	LEU	CB	-42.00	-21.36	55.08	15.00
195	LEU	CG		-20.66	56.16	15.00
195	LEU	CD1	-44.27	-20.65	55.77	15.00
195	LEU	CD2	-42.60	-21.36	57.49	15.00
195	LEU	C	-39.83	-22.09	54.09	15.00
195	LEU	0	-39.85	-21.47	53.04	15.00
196	MET	N	-39.17	-23.23	54.23	15.00

#### · TABLE V

196 MET CA	-38.49	-23.84	53.10	15.00
196 MET CB	-37.01	-24.08	53.39	15.00
196 MET CG	-36.15	-22.83	53.37	15.00
196 MET SD	-34.45	-23.19	53.93	15.00
196 MET CE	-33.63	-23.67	52.36	15.00
196 MET C	-39.17	-25.14	52.72	15.00
196 MET 0	-39.59	-25.89	53.59	15.00
197 ALA N	-39.22	-25.41	51.41	15.00
197 ALA CA	-39.86	-26.62	50.87	15.00
197 ALA CB	-39.64	-26.70	49.36	15.00
197 ALA C	-39.42	-27.93	51.53	15.00
197 ALA O	-38.23	-28.21	51.67	15.00
198 ARG N	-40.41	-28.73	51.91	15.00
198 ARG CA	-40.18	-30.01	52.57	15.00
198 ARG CB	-40.77	-30.03	53.98	15.00
198 ARG CG	-40.78	-31.39	54.66	15.00
198 ARG CD	-41.18	-31.28	56.12	15.00
198 ARG NE	-42.52	-30.73	56.31	15.00
198 ARG CZ	-43.63	-31.47	56.40	15.00
198 ARG NH1	-43.55	-32.80	56.31	15.00
198 ARG NH2	-44.80	-30.89	56.62	15.00
198 ARG C	-40.74	-31.13	51.71	15.00
198 ARG O	-41.84	-31.00	51.16	15.00
199 ASN N	-39.98	-32.21	51.61	15.00
199 ASN CA	-40.35	-33.37	50.81	15.00
199 ASN CB	-41.72	-33.92	51.23	15.00
199 ASN CG	-41.71	-34.55	52.61	15.00
199 ASN OD1	-40.67	-34.60	53.26	15.00
199 ASN ND2	-42.87	-35.01	53.07	15.00
199 ASN C	-40.31	-33.04	49.32	15.00
199 ASN 0	-41.18	-33.43	48.57	15.00
200 LYS N	-39.30	-32.27	48.92	15.00
200 LYS CA	-39.13		47.54	15.00
200 LYS CB	-39.46	-30.41	47.32	15.00
200 LYS CG	-39.74	-30.07	45.87	15.00
200 LYS CD	-41.24	-30.05	45.59	15.00
200 LYS CE	-41.92	-28.93	46.40	15.00
200 LYS NZ		-28.84	46.21	15.00
200 LYS C	-37.68			15.00
200 LYS 0	-36.89	-31.26	46.91	15.00
201 ASN N	-37.34	-33.46	47.14	15.00
201 ASN CA	-36.00	-33.95	46.83	15.00

201 ASN CB	-25 70	34.00		
201 ASN CG	-35.78 -36.19			
201 ASN OD1		_	44.59	15.00
201 ASN ND2	-35.40		43.87	
201 ASN C	-34.84		44.76	
201 ASN 0	-33.84			
202 ASN N	-34.98		46.92	15.00
202 ASN CA	-33.97		48.86	15.00
202 ASN CB	-32.74		_	15.00
202 ASN CG	-31.91	· - <del>-</del>	-	15.00
202 ASN OD1	-32.36		51.13	15.00
202 ASN ND2	-30.67		52.07	15.00
202 ASN C	-33.56		51.13	15.00
202 ASN 0	-32.39	-31.08	49.33	15.00
203 ALA N	-34.51	-30.82	49.02	15.00
203 ALA H	-35.31	-30.16 -30.45	49.36	15.00
203 ALA CA	-34.34	-30.45	49.85	15.00
203 ALA CB	-35.57		48.93	15.00
203 ALA C	-33.15	-28.14	49.27	15.00
203 ALA O		-28.14 -27.98	49.67	15.00
204 CYS N	-32.16		50.89	15.00
204 CYS CA	-30.95	-27.06	48.86	15.00
204 CYS C	-30.08	-27.85	49.31 50.28	15.00
204 CYS 0	-29.25	-27.26	50.28	15.00
204 CYS CB	-31.27	-25.68	49.90	15.00
204 CYS SG	-32.21	-24.52	48.84	15.00 15.00
205 GLY N	-30.24	-29.17	50.32	15.00
205 GLY CA	-29.45	-29.99	51.22	15.00
205 GLY C	-29.93	-29.89	52.66	15.00
205 GLY O	-29.14	-30.07	53.60	15.00
206 ILE N	-31.23	-29.68	52.83	15.00
206 ILE CA	-31.84	-29.52	54.15	15.00
206 ILE CB	-33.39	-29.24	54.01	15.00
206 ILE CG2		-30.44	53.42	15.00
206 ILE CG1		-28.84	55.35	15.00
206 ILE CD1	-33.66	-27.44	55.81	15.00
206 ILE C	-31.57	-30.69	55.11	15.00
206 ILE O		-30.47	56.31	15.00
207 ALA N	-31.46	-31.91	54.59	15.00
207 ALA CA	-31.21	-33.09	55.42	15.00
207 ALA CB		-34.12	55.21	15.00
207 ALA C	-29.83	-33.73	55.24	15.00
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207 ALA O	-29.58	-34.86	55.66	15.00
208 ASN N	-28.92	-32.98	54.65	
208 ASN CA	-27.58		54.42	15.00
208 ASN CB	-26.91	-32.79	53.25	15.00
208 ASN CG	-27.23	-33.43	51.92	15.00
208 ASN OD1	-27.61	-34.59		15.00
208 ASN ND2	-27.06	-32.67	50.84	15.00
208 ASN C	-26.68	-33.44	55.65	15.00
208 ASN O	-25.80	-34.29	55.81	15.00
209 LEU N	-26.91	-32.46	56.52	15.00
209 LEU CA	-26.09	-32.29	57.72	15.00
209 LEU CB	-24.96	-31.30	57.44	15.00
209 LEU CG	-23.70		58.30	15.00
209 LEU CD1	-22.81	-32.53	57.75	15.00
209 LEU CD2	-22.95	-30.12	58.33	15.00
209 LEU C	-26.86	-31.91	58.99	15.00
209 LEU O	-26.42	-31.04	59.74	15.00
210 ALA N	-27.99	-32.56	59.23	15.00
210 ALA H	-27.89	-32.43	58.64	15.00
210 ALA CA	-28.78	-32.25	60.42	15.00
210 ALA CB	-29.43	-31.67	60.51	15.00
210 ALA C	-28.24	-33.03	61.63	15.00
210 ALA O	-27.87	-34.20	61.51	15.00
211 SER N	-28.17	-32.35	62.78	15.00
211 SER CA	-27.71	-32.98	64.03	15.00
211 SER CB	-26.19	-33.01	64.15	15.00
211 SER OG	-25.65	-31.71	64.29	15.00
211 SER C	-28.32	-32.28	65.24	15.00
211 SER O	-28.93	-31.20	65.11	15.00
212 PHE N	-28.18	-32.89	66.41	15.00
212 PHE CA	-28.74	-32.30	67.61	15.00
212 PHE CB	-30.25	-32.56	67.71	15.00
212 PHE CG		-34.00	67.79	15.00
212 PHE CD1	-31.04	-34.69	66.66	15.00
212 PHE CD2	-30.53	-34.59	69.00	15.00
212 PHE CE1	-31.36	-36.03	66.73	15.00
212 PHE CE2	-30.85	-36.04	69.07	15.00
212 PHE CZ		-36.71	67.94	15.00
212 PHE C		-32.78	68.87	15.00
212 PHE 0	-27.55	-33.94	68.95	15.00
213 PRO N		-31.91	69.88	15.00
213 PRO CD		-30.60	69.99	15.00
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213 PRO CA		-32.25		15.00
213 PRO CB	-27.11		71.80	15.00
213 PRO CG	-28.44	-30.26	71.47	15.00
213 PRO C	-28.13	-33.17	71.96	15.00
213 PRO O	-29.36	-33.23	71.79	15.00
214 LYS N	-27.48	-33.94	72.82	15.00
214 LYS CA	-28.19	-34.82	73.72	15.00
214 LYS CB	-27.71	-36.27	73.59	15.00
214 LYS CG	-27.91	-36.90	72.23	15.00
214 LYS CD	-27.28	-38.29	72.16	15.00
214 LYS CE	-25.79	-38.27	72.50	15.00
214 LYS NZ	-25.11	-39.56	72.18	15.00
214 LYS C	-27.84	-34.26	75.07	15.00
214 LYS 0	-26.76	-33.71	75.24	15.00
215 MET N	-28.79	-34.27	75.99	15.00
215 MET CA	-28.55	-33.78	77.34	15.00
215 MET CB	-29.47	-32.60	77.70	15.00
215 MET CG	-29.12	-31.30	76.96	15.00
215 MET SD	-29.59	-29.73	77.80	15.00
215 MET CE	-28.68	-28.54	76.81	15.00
215 MET C	-28.66	-34.93	78.34	15.00
215 MET OT1	-27.81	-35.01	79.26	15.00
215 MET OT2	-29.54	-35.80	78.16	15.00
216 нон он2	-40.32	-20.86	90.40	15.00
217 нон он2	-20.71	-32.43	79.67	15.00
218 нон он2	-31.33	-16.38	65.47	15.00
219 НОН ОН2	-29.76	-17.63	70.42	15.00
220 нон он2	-7.13	-18.39	66.48	15.00
221 НОН ОН2	-15.45	-12.55	73.01	15.00
222 НОН ОН2	-34.69	-23.23	69.94	15.00
223 НОН ОН2	-11.03	-30.64	72.74	15.00
224 НОН ОН2	-30.92	-18.33	68.20	15.00
225 НОН ОН2	-24.49	-30.79	61.59	15.00
226 НОН ОН2		-10.95		15.00
227. НОН ОН2		-21.84		15.00
228 НОН ОН2		-29.94	49.49	15.00
229 НОН ОН2		-34.98	55.75	15.00
230 НОН ОН2		-12.36	67.54	15.00
231 HOH OH2			60.21	15.00
232 НОН ОН2		-33.51		15.00
233 HOH OH2		-25.41	63.34	15.00
234 нон он2	-57.83	-31.91	39.28	15.00

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235 HOH OH2		-20.10	63.45	15.00
236 НОН ОН2				
237 НОН ОН2			68.27	
238 НОН ОН2	-28.19			
239 НОН ОН2	-26.07			
240 нон он2	-35.84	-27.16	51.26	
241 HOH OH2		-24.80	49.57	
242 HOH OH2		-32.65		15.00
243 HOH OH2		-9.00	77.82	15.00
244 HOH OH2		-14.85	64.38	15.00
245 HOH OH2	-18.39	-3.01	63.15	
246 нон он2	~33.49		70.47	
247 НОН ОН2	-48.24		79.27	15.00
248 HOH OH2	-17.16	-11.08	74.86	15.00
249 НОН ОН2	-7.77	-18.99	72.85	15.00
250 нон он2	-12.50	-24.63	81.88	15.00
251 нон он2	-28.11		58.10	15.00
252 НОН ОН2	-35.24		53.39	15.00
253 нон он2	-31.85	-28.95	46.18	15.00
254 нон он2	-35.11	-24.97	46.75	15.00
255 нон он2	-42.46	-38.44	54.37	15.00
256 нон он2	-37.82	-16.40	67.58	15.00
257 НОН ОН2	-43.11	-16.23		15.00
258 нон он2	-36.79		73.69	15.00
259 нон он2	-34.92	-15.40	75.95	15.00
260 нон он2	-32.03	-7.39	60.30	15.00
261 HOH OH2	-19.94	-8.07	62.81	15.00
262 HOH OH2	-33.79	-20.76	69.68	15.00
263 нон он2	-33.86		74.42	15.00
264 HOH OH2	-11.97	-27.02	71.08	15.00
265 HOH OH2	-8.26	-25.33	61.28	15.00
266 нон он2	-19.53	-42.28	58.81	15.00
267 НОН ОН2	-20.68	-32.75		15.00
268 нон он2	-24.87	-33.89	60.62	15.00
269 НОН ОН2	-2.83	-32.79	71.85	15.00
270 нон он2	-14.43	-40.52	59.53	15.00
271 нон он2	-21.46	-37.41	78.35	15.00
272 НОН ОН2		-36.03	71.33	15.00
273 нон он2	-28.57	-35.40	88.70	15.00
274 НОН ОН2	-13.04	-12.02	63.26	15.00
275 НОН ОН2		-11.89	72.80	15.00
276 нон он2		-30.13	56.41	15.00

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277	HOH	OH2	-29.86	-20.69	48.27	15.00
278	HOH	OH2	-26.77	-22.94	44.37	15.00
279	HOH	OH2	-25.17	-36.24	49.68	15.00
280	HOH	OH2	-19.40	-31.57	49.99	15.00
281	HOH	OH2	-34.95	-29.42	45.52	15.00
282	HOH	OH2	-37.69	-30.43	50.51	15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors  $(\mathbb{A}^2)$  for the cathepsin K complex with inhibitor 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)] carbohydrazide.

Residue Atom	x	Y	z	В
1 ALA CB	-53.28	-28.69	64.46	15.00
1 ALA C	-53.74	-30.77	63.13	15.00
1 ALA O	-54.17	-31.71	63.79	15.00
1 ALA N	-55.61	-29.36	63.92	15.00
1 ALA CA	-54.20	-29.34	63.43	15.00
2 PRO N	-52.92	-30.93	62.07	15.00
2 PRO CD	-52.55	-29.87	61.11	15.00
2 PRO CA	-52.38	-32.23	61.65	15.00
2 PRO CB	-52.22	-32.03	60.15	15.00
2 PRO CG	-51.68	-30.61	60.09	15.00
2 PRO C	-51.02	-32.37	62.31	15.00
2 PRO O	-50.88	-32.09	63.50	15.00
3 ASP N	-50.02	-32.75	61.52	15.00
3 ASP CA	-48.67	-32.92	62.02	15.00
3 ASP CB	-47.96	-34.03	61.25	15.00
3 ASP CG	-48.48	-35.41	61.59	15.00
3 ASP OD1	-49.68	~35.69	61.38	15.00
3 ASP OD2	-47.66	-36.24	62.06	15.00
3 ASP C	-47.93	-31.60	61.84	15.00
3 ASP O	-47.35	-31.34	60.78	15.00
4 SER N	-48.02	-30.74	62.84	15.00
4 SER CA	-47.34	-29.45	62.82	15.00
4 SER CB	-48.32	-28.34	62.42	15.00
4 SER OG	-48.91	-28.65	61.17	15.00
4 SER C	-46.76	-29.17	64.20	15.00
4 SER O	-47.33	-29.58	65.22	15.00
5 VAL N	<b>-45.6</b> 0	-28.54	64.23	15.00
5 VAL CA	-45.00	-28.20	65.51	15.00
5 VAL CB	-44.16	-29.36	66.11	15.00
5 VAL CG1	-42.89	-29.57	65.35	15.00
5 VAL CG2	-43.87	-29.08	67.57	15.00
5 VAL C	-44.21	-26.91	65.37	15.00
5 VAL O	-43.46	-26.73	64.41	15.00
6 ASP N	-44.48	-25.98	66.27	15.00

	ARC		-26.86	-36.86	79.28	15.00
108	3 ARG	NE	-26.63	-38.14	78.60	15.00
108	ARC	CZ	-27.58	-38.89	78.03	15.00
108	ARG	NH1	-28.85	-38.51	78.05	15.00
108	ARG	NH2	-27.24	-40.03	77.45	15.00
108	ARG	C	-23.64	-35.78	76.18	15.00
108	ARG	0	-23.84	-36.99	76.06	15.00
109	GLY	N	-23.20	-35.03	75.17	15.00
109	GLY	CA	-22.97	-35.63	73.87	15.00
109	GLY	C	-23.82	-34.99	72.80	15.00
109	GLY	0	-24.27	-33.85	72.95	15.00
110	TYR	N	-24.09	-35.72	71.72	15.00
110		CA	-24.89	-35.19	70.63	15.00
110	TYR	CB	-24.12	-34.08	69.91	15.00
	TYR		-22.86	-34.56	69.20	15.00
110	TYR	CD1	-21.64	-34.62	69.87	15.00
110	TYR	CE1	-20.49	-35.03	69.22	15.00
110		CD2	-22.90	-34.93	67.85	15.00
110	TYR	CE2	-21.75	-35.34	67.19	15.00
110	TYR	CZ	-20.55	-35.39	67.88	15.00
110		OH	-19.39	-35.77	67.23	15.00
110		С	-25.22	-36.28	69.62	15.00
110	TYR	0	-24.44	-37.21	69.43	15.00
	ARG		-26.34	-36.11	68.92	15.00
	ARG		-26.76	-37.08	67.92	15.00
	ARG		-28.02	-37.81	68.38	15.00
	ARG		-27.81	-38.84	69.48	15.00
	ARG		-28.75	-40.03	69.33	15.00
	ARG		-28.01	-41.28	69.36	15.00
	ARG		-27.89	-42.06	70.44	15.00
	ARG		-27.18	-43.18	70.36	15.00
	ARG		-28.50	-41.73	71.58	15.00
	ARG		-26.99	-36.46	66.54	15.00
111	ARG		-27.67	-35.45	66.41	15.00
112			-26.41	-37.06	65.50	15.00
112			-26.56	-36.57	64.14	15.00
112	GLU			-37.00	63.29	15.00
112	GLU			-36.42	63.76	15.00
112				-35.56	62.70	15.00
	GLU			-34.57	62.22	15.00
	GLU			-35.88	62.33	15.00
112	GLU			-37.13	63.56	15.00
112	GLU	C	-28.64	-37.75	64.27	15.00

113 ILE N	-28.13		62.29	15.00
113 ILE CA	-29.35		61.64	15.00
113 ILE CB	-30.38		61.34	
113 ILE CG2	-31.67	-36.83	60.78	
113 ILE CG1	-30.71	-35.45	62.60	15.00
113 ILE CD1	-31.50	-36.20	63.65	15.00
113 ILE C	-28.85	-37.98	60.33	15.00
113 ILE 0	-27.91	-37.46	59.73	15.00
114 PRO N	-29.41	-39.13	59.92	15.00
114 PRO CD	-30.48	-39.91	60.57	15.00
114 PRO CA	-28.98	-39.77	58.68	15.00
114 PRO CB	-30.10	-40.78	58.43	15.00
114 PRO CG	-30.41	-41.23		15.00
114 PRO C	-28.87	-38.79	57.54	15.00
114 PRO O	-29.84	-38.12	57.20	15.00
115 GLU N	-27.67	-38.66	56.99	15.00
115 GLU CA	-27.46	-37.74	55.89	15.00
115 GLU CB	-26.07		55.28	15.00
115 GLU CG	-24.92	-37.19	56.01	15.00
115 GLU CD	-23.60	-37.16	55.20	15.00
115 GLU OE1	-23.65	-37.20	53.94	15.00
115 GLU OE2	-22.51	-37.08	55.82	15.00
115 GLU C	-28.53	-37.91	54.82	15.00
115 GLU O	-28.70	-39.00	54.27	15.00
116 GLY N	-29.27	-36.83	54.59	15.00
116 GLY CA	-30.30	-36.78	53.57	15.00
116 GLY C	-31.63	-37.36	53.99	15.00
116 GLY O	-32.46	-37.66	53.13	15.00
117 ASN N	-31.88	-37.44	55.29	15.00
117 ASN CA	-33.12	-38.03	55.77	15.00
117 ASN CB	-32.80	-39.09	56.82	15.00
117 ASN CG	-33.80	-40.21	56.84	15.00
117 ASN OD1	-35.02	-39.99	56.90	15.00
117 ASN ND2	-33.31	-41.44	56.76	15.00
117 ASN C	-34.17	-37.06	56.31	15.00
117 ASN 0	-34.17	-36.72	57.50	15.00
118 GLU N	-35.12	-36.66	55.48	15.00
118 GLU CA	-36.15	-35.72	55.91	15.00
118 GLU CB	-36.86	-35.08	54.72	15.00
118 GLU CG	-36.09	-33.98	54.02	15.00
118 GLU CD	-36.99	-33.14	53.15	15.00
118 GLU OE1	-36.58	-32.75	52.04	15.00
118 GLU OE2	-38.13	-32.87	53.57	15.00

118 GLU C	-37.18	-36.33	56.83	15.00
118 GLU O	-37.76			15.00
119 LYS N	-37.46	-37.61	56.66	15.00
119 LYS CA	-38.43	~38.29	57.53	15.00
119 LYS CB	-38.66	-39.71	57.03	15.00
119 LYS CG	-39.20		55.60	15.00
119 LYS CD	-40.54	-39.04	55.49	15.00
119 LYS CE	-41.04	-38.97	54.07	15.00
119 LYS NZ	-40.26	-38.00	53.26	15.00
119 LYS C	-37.89	-38.30	58.95	15.00
119 LYS O	-38.55	-37.86	59.89	15.00
120 ALA N	-36.64	-38.74	59.07	15.00
120 ALA CA	-35.94	-38.81	60.34	15.00
120 ALA CB	-34.58	-39.42	60.15	15.00
120 ALA C	-35.81	-37.43	60.97	15.00
120 ALA O	-35.88	-37.28	62.19	15.00
121 LEU N	-35.56	-36.42	60.14	15.00
121 LEU CA	-35.43	-35.06	60.66	15.00
121 LEU CB	-34.96	-34.10	59.56	15.00
121 LEU CG	-34.93	-32.61	59.96	15.00
121 LEU CD1	-33.81	-32.34	60.93	15.00
121 LEU CD2	-34.78	-31.76	58.72	15.00
121 LEU C	-36.76	-34.60	61.25	15.00
121 LEU O	-36.81	-33.98	62.31	15.00
122 LYS N	-37.86	-34.91	60.57	15.00
122 LYS CA	-39.17	-34.53	61.06	15.00
122 LYS CB	-40.26	-35.04	60.11	15.00
122 LYS CG	-41.65	-34.70	60.59	15.00
122 LYS CD	-42.73	-35.28	59.72	15.00
122 LYS CE	-44.10	-34.79	60.19	15.00
122 LYS NZ	-45.21	-35.17	59.28	15.00
122 LYS C	-39.36	-35.15	62.43	15.00
122 LYS 0	-39.79	-34.47	63.36	15.00
123 ARG N	-39.02	-36.44	62.55	15.00
123 ARG CA		-37.17	63.81	15.00
123 ARG CB	-38.72	-38.64	63.64	15.00
123 ARG CG	-39.86	-39.56	63.26	15.00
123 ARG CD	-39.43	-41.02	63.33	15.00
123 ARG NE		-41.48	62.14	15.00
123 ARG CZ		-41.79		15.00
123 ARG NH1		-41.70	63.21	15.00
123 ARG NH2		-42.22	60.99	15.00
123 ARG C	-38.28	-36.55	64.90	15.00

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123 ARG O	-38.69	-36.51	66.05	15.00
124 ALA N	-37.09	,		
124 ALA CA	-36.20	-35.45		
124 ALA CB	-34.85	-35.18		
124 ALA C	-36.81			
124 ALA O	-36.96			
125 VAL N	-37.17			15.00
125 VAL CA	-37.77	-31.97		15.00
125 VAL CB	-38.07	-31.11	64.25	15.00
125 VAL CG1	-38.97	-29.94	64.59	15.00
125 VAL CG2	-36.78			
125 VAL C	-39.04			15.00
125 VAL 0	-39.36	-31.43	67.20	15.00
126 ALA N	-39.74		66.00	15.00
126 ALA CA	-40.97		66.69	15.00
126 ALA CB	-41.78		65.88	15.00
126 ALA C	-40.72		68.09	15.00
126 ALA O	-41.37		69.06	15.00
127 ARG N		-35.10	68.19	15.00
127 ARG CA	-39.41	-35.74	69.44	15.00
127 ARG CB	-38.71	-37.07	69.18	15.00
127 ARG CG	-39.58	-38.30	69.27	15.00
127 ARG CD	-40.17	-38.70	67.94	15.00
127 ARG NE	-39.87	-40.09	67.60	15.00
127 ARG CZ	-40.52	-40.78	66.67	15.00
127 ARG NH1	-40.18	-42.03	66.40	15.00
127 ARG NH2	-41.55	-40.25	66.03	15.00
127 ARG C	-38.52	-34.91	70.35	15.00
127 ARG 0	-38.86	-34.71	71.52	15.00
128 VAL N	-37.37	-34.50	69.82	15.00
128 VAL CA	-36.36	-33.72	70.56	15.00
128 VAL CB	-34.95	-33.91	69.94	15.00
128 VAL CG1	-33.88	-33.45	70.91	15.00
128 VAL CG2	-34.73	-35.35	69.57	15.00
128 VAL C		-32.22	70.66	15.00
128 VAL 0		-31.61	71.71	15.00
129 GLY N		-31.60	69.55	15.00
129 GLY CA		-30.17	69.54	
129 GLY C		-29.54	68.53	15.00
129 GLY O		-30.22	67.58	15.00
130 PRO N		-28.25	68.71	15.00
130 PRO CD		-27.38		15.00
130 PRO CA		-27.50	69.78	15.00
	JJ. UM	27.30	67.83	15.00

130 P	RO CB	-34.80	-26.22	68.62	15.00
130 PI	RO CG	-36.12		69.26	15.00
130 PI	RO C	-33.74	-28.21	67.50	15.00
130 P	RO O	-32.90	-28.46	68.37	15.00
131 V	AT N	-33.59	-28.51	66.21	15.00
131 V	AL CA	-32.42	-29.19	65.68	15.00
131 V	T CB	-32.84	-30.34	64.73	15.00
131 V	L CG1	-31.64	-31.06	64.18	15.00
131 VA	L CG2	-33.74	-31.33	65.48	15.00
131 VZ	ТC	-31.57	-28.19	64.90	15.00
131 VA	T O	-32.08	-27.25	64.28	15.00
132 SE	R N	-30.25	-28.38	64.96	15.00
132 SE	R CA	-29.32	-27.53	64.26	15.00
132 SE	R CB	-28.00	-27.48	65.02	15.00
132 SE	R OG	-28.18	-26.93	66.31	15.00
132 SE	R C	-29.08	-28.14	62.88	15.00
132 SE	R O	-28.67	-29.30	62.79	15.00
133 VA	LN	-29.38	-27.38	61.83	15.00
133 VA	L CA	-29.21	-27.83	60.44	15.00
133 VA	L CB	-30.57	-27.93	59.68	15.00
133 VA	L CG1	-31.55	-28.81	60.43	15.00
133 VA	L CG2	-31.16	-26.56	59.49	15.00
133 VA	r c	-28.30	-26.86	59.68	15.00
133 VA	LO	-28.06	-25.74	60.13	15.00
134 AL	A N	-27.79	-27.28	58.52	15.00
134 AL	A CA	-26.94	-26.44	57.69	15.00
134 AL	A CB	-25.50	-26.93	57.73	15.00
134 AL	A C	-27.49	-26.50	56.28	15.00
13 <b>4 AL</b>	A O	-27.84	-27.57	55.79	15.00
135 IL	E N	-27.58	-25.36	55.61	15.00
135 IL	E CA	-28.14	-25.32	54.27	15.00
135 IL	E CB	-29.58	-24.73	54.30	15.00
	E CG2	-30.49	-25.57	55.19	15.00
135 IL	E CG1	-29.53	-23.26	54.76	15.00
135 IL	E CD1	-30.88	-22.58	54.88	15.00
135 IL	EC	-27.33	-24.44	53.32	15.00
135 IL	E O	-26.33	-23.82	53.72	15.00
136 AS	PN	-27.75	-24.41	52.05	15.00
136 AS	P CA	-27.12	-23.57	51.05	15.00
136 AS	P CB	-27.25	-24.16	49.65	15.00
136 AS	P CG	-26.77	-23.20	48.57	15.00
136 ASI	P OD1	-27.27	-23.28	47.43	15.00
136 AS	P OD2	-25.90	-22.35	48.85	15.00

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136 ASP C	-27.87	-22.24	51.10	15.00
136 ASP 0	-29.05	-22.17	50.77	15.00
137 ALA N	-27.19	-21.21	51.59	15.00
137 ALA CA	-27.80	-19.89	51.68	15.00
137 ALA CB	-27.61	-19.32	53.08	15.00
137 ALA C	-27.23	-18.94	50.62	15.00
137 ALA O	-27.64	-17.78	50.53	15.00
138 SER N	-26.26	-19.43	49.84	15.00
138 SER CA	-25.66	-18.65	48.77	15.00
138 SER CB	-24.30	-19.22	48.37	15.00
138 SER OG	-23.28	-18.86	49.30	15.00
138 SER C	-26.61	-18.73	47.60	15.00
138 SER O	-26.51	-19.65	46.80	15.00
139 LEU N	-27.54	-17.77	47.57	15.00
139 LEU CA	-28.59	-17.66	46.55	15.00
139 LEU CB	-29.53	-18.86	46.65	15.00
139 LEU CG	-29.89	-19.63	45.38	15.00
139 LEU CD1	-28.66	-20.27	44.80	15.00
139 LEU CD2	-30.92	-20.68	45.71	15.00
139 LEU C	-29.39	-16.39	46.85	15.00
139 LEU O	-29.91	-16.22	47.96	15.00
140 THR N	-29.52	-15.49	45.87	15.00
140 THR CA	-30.23	-14.23	46.08	15.00
140 THR CB	-30.06	-13.28	44.87	15.00
140 THR OG1	-29.05	-13.81	43.99	15.00
140 THR CG2	-29.62	-11.89	45.34	15.00
140 THR C	-31.71	-14.37	46.42	15.00
140 THR 0	-32.23	-13.63	47.24	15.00
141 SER N	-32.39	-15.35	45.83	15.00
141 SER CA	-33.80	-15.59	46.10	15.00
141 SER CB	-34.31	-16.76	45.27	15.00
141 SER OG	-33.42	-17.86	45.34	15.00
141 SER C	-34.00	-15.89	47.57	15.00
141 SER O	-35.03		48.16	15.00
142 PHE N	-33.01	-16.51	48.18	15.00
142 PHE CA	-33.07	-16.86	49.59	15.00
142 PHE CB		-17.88	49.93	15.00
142 PHE CG		-18.24	51.39	15.00
142 PHE CD1		-19.25	51.89	15.00
142 PHE CD2		-17.54	52.25	15.00
142 PHE CE1		-19.58	53.23	15.00
142 PHE CE2		-17.85	53.60	15.00
L42 PHE CZ	-31.85	-18.87	54.09	15.00

142 PHE C	-32.88	-15.61	50.43	15.00
142 PHE 0	-33.66	-15.33		15.00
143 GLN N	-31.86	-14.84	50.09	15.00
143 GLN CA	-31.54	-13.64	50.83	15.00
143 GLN CB	-30.25	-13.06	50.30	15.00
143 GLN CG	-29.18	-14.11	50.24	15.00
143 GLN CD	-27.84	-13.53	49.97	15.00
143 GLN OE1	-27.41	-12.58	50.62	15.00
143 GLN NE2	-27.15	-14.10	48.99	15.00
143 GLN C	-32.63	-12.59	50.89	15.00
143 GLN 0	-32.90	-12.05	51.95	15.00
144 PHE N	-33.29	-12.29	49.77	15.00
144 PHE CA	-34.36	-11.28	49.80	15.00
144 PHE CB	-34.33	-10.38	48.55	15.00
144 PHE CG	-34.46	-11.12	47.24	15.00
144 PHE CD1	-33.45	-11.04	46.30	15.00
144 PHE CD2	-35.59	-11.87	46.95	15.00
144 PHE CE1	-33.56	-11.71	45.09	15.00
144 PHE CE2	-35.71	-12.55	45.75	15.00
144 PHE CZ	-34.70	-12.46	44.82	15.00
144 PHE C	-35.76	-11.83	50.04	15.00
144 PHE 0	-36.76	-11.19	49.68	15.00
145 TYR N	-35.83	-13.01	50.66	15.00
145 TYR CA	-37.09	-13.68	50.97	15.00
145 TYR CB	-36.80	-15.04	51.62	15.00
145 TYR CG	-37.95	-15.65	52.36	15.00
145 TYR CD1	-38.83	-16.52	51.73	15.00
145 TYR CE1	-39.93	-17.05	52.41	15.00
145 TYR CD2	-38.20	-15.31	53.69	15.00
145 TYR CE2	-39.28	-15.83	54.38	15.00
145 TYR CZ	-40.14	-16.69	53.73	15.00
145 TYR OH	-41.24	-17.16	54.42	15.00
145 TYR C	-37.87	-12.81	51.93	15.00
145 TYR O	-37.27	-12.10	52.74	15.00
146 SER N	-39.20	-12.88	51.89	15.00
146 SER CA	-40.01	-12.07	52.78	15.00
146 SER CB	-40.45	-10.78	52.10	15.00
146 SER OG	-41.31	-11.05	51.01	15.00
146 SER C	-41.23	-12.81	53.34	15.00
46 SER O	-41.69	-12.52	54.45	15.00
47 LYS N	-41.77	-13.75	52.58	15.00
L47 LYS CA		-14.52	53.04	15.00
147 LYS CB	-44.16	-13.62	53.21	15.00

147 LYS CG	-44.84	-13.14	51.92	15.00
147 LYS CD	-45.64	-11.86	52.19	15.00
147 LYS CE	-44.71	-10.72	52.70	15.00
147 LYS NZ	-45.41	-9.52	53.28	15.00
147 LYS C	-43.27	-15.68	52.11	15.00
147 LYS O	-43.03	-15.62	50.91	15.00
148 GLY N	-43.83	-16.74	52.69	15.00
148 GLY CA	-44.21	-17.90	51.90	15.00
148 GLY C	-43.31	-19.08	52.16	15.00
148 GLY O	-42.50	-19.07	53.08	15.00
149 VAL N	-43.46	-20.12	51.35	15.00
149 VAL CA	-42.65	-21.32	51.50	15.00
149 VAL CB	-43.52	-22.60	51.39	15.00
149 VAL CG1	-42.66	-23.84	51.53	15.00
149 VAL CG2	-44.59	-22.59	52.45	15.00
149 VAL C	-41.57	-21.27	50.42	15.00
149 VAL 0	-41.84	-21.44	49.24	15.00
150 TYR N	-40.34	-21.01	50.84	15.00
150 TYR CA	-39.21	-20.90	49.93	15.00
150 TYR CB	-37.98	-20.37	50.67	15.00
150 TYR CG	-36.75	-20.27	49.80	15.00
150 TYR CD1	-36.73	-19.44	48.68	15.00
150 TYR CE1	-35.61	-19.36	47.86	15.00
150 TYR CD2	-35.61	-21.03	50.09	15.00
150 TYR CE2 150 TYR CZ	-34.48	-20.96	49.27	15.00
	-34.49	-20.13	48.16	15.00
150 TYR OH 150 TYR C	-33.41	-20.06	47.32	15.00
150 TYR O	-38.81	-22.16	49.19	15.00
150 TYR N	-38.56	-23.19	49.80	15.00
151 TYR CA	-38.68 -38.24	-22.03	47.88	15.00
151 TYR CB		-23.13	47.02	15.00
151 TYR CG	-39.38 -38.89	-24.11	46.72	15.00
151 TYR CD1		-25.32 -26.01	45.96	15.00
151 TYR CE1	-37.74 -37.23	-20.01	46.37	15.00
151 TYR CD2			45.63	15.00
151 TYR CE2		-25.74	44.79	15.00
151 TYR CZ		-26.80 -27.45	44.03	15.00
151 TYR OH			44.46	15.00
L51 TYR C		-28.50 -22.50	43.72	15.00
L51 TYR O		-22.58	45.73	15.00
L52 ASP N		-21.69 -23.12	45.08	15.00
152 ASP CA		-23.12 -22.69	45.37	15.00
CA	-33.14	-44.69	44.19	15.00

152	ASP	CB	-34.80	-21.56	44.59	15.00
152	ASP	CG	-34.09	-20.92	43.42	15.00
152	ASP	OD1	-33.27	-21.60	42.76	15.00
152	ASP	OD2	-34.35	-19.73	43.17	15.00
152	ASP	С	-34.93	-23.88	43.70	15.00
152	ASP	0	-33.98	-24.30	44.37	15.00
153	GLU	N	-35.23	-24.36	42.51	15.00
153	GLU	CA	-34.55	-25.53	41.95	15.00
153	GLU	CB	-35.15	-25.91	40.59	15.00
153	GLU	CG	-35.40	-24.74	39.62	15.00
153	GLU	CD	-34.12	-24.02	39.16	15.00
153	GLU	OE1	-33.16	-24.68	38.70	15.00
153	GLU	OE2	-34.09	-22.76	39.27	15.00
153	GLU	С	-33.03	-25.50	41.88	15.00
153	GLU	0	-32.40	-26.49	41.49	15.00
154	SER	N	-32.42	-24.37	42.22	15.00
154	SER	CA	-30.96	-24.26	42.18	15.00
154	SER	CB	-30.53	-23.00	41.43	15.00
154	SER	OG	-30.93	-23.09	40.06	15.00
154	SER	C	-30.27	-24.34	43.55	15.00
154	SER	0	-29.04	-24.33	43.65	15.00
155	CYS	N	-31.06	-24.43	44.61	15.00
155	CYS	CA	-30.49	-24.54	45.95	15.00
155	CYS	С	-29.66	-25.81	45.88	15.00
155	CYS	0	-30.16	-26.88	45.54	15.00
155	CYS	CB	-31.60	-24.69	46.96	15.00
155	CYS	SG	-31.20	-23.89	48.53	15.00
156	ASN	N	-28.36	-25.70	46.16	15.00
156	ASN	CA	-27.50	-26.87	46.05	15.00
156	ASN	CB	-26.20	-26.49	45.36	15.00
156	asn	CG	-25.34	-27.69	45.04	15.00
156	ASN	OD1	-25.83	-28.82	44.97	15.00
156	asn	ND2	-24.05	-27.47	44.88	15.00
156	ASN	С	-27.21	-27.66	47.33	15.00
156	ASN	0	-26.37	-27.26	48.13	15.00
157	SER	N	-27.82	-28.83	47.43	15.00
157	SER	CA	-27.66	-29.72	48.57	15.00
157	SER	CB	-28.41	-31.03	48.33	15.00
157	SER	OG	-29.78	-30.78	48.03	15.00
157	SER	С	-26.19	-30.04	48.87	15.00
157	SER	0	-25.86	-30.54	49.96	15.00
158	ASP	N		-29.75	47.92	15.00
158	ASP	CA	-23.89	-30.03	48.07	15.00

158 ASP CB	-23.30	-30.43	46.71	15.00
158 ASP CG	-24.01			15.00
158 ASP OD1	-25.23	-31.53	45.79	15.00
158 ASP OD2	-23.34	-32.67		15.00
158 ASP C	-23.11	-28.86	48.64	15.00
158 ASP O	-22.00	-29.02	49.15	15.00
159 ASN N	-23.68	-27.66	48.60	15.00
159 ASN CA	-22.98	-26.49	49.11	15.00
159 ASN CB	-23.02	-25.36		15.00
159 ASN CG	-21.95	-24.29	48.31	15.00
159 ASN OD1	-21.61	-23.96	49.45	15.00
159 ASN ND2	-21.43	-23.72	47.22	15.00
159 ASN C	-23.59	-26.01	50.41	15.00
159 ASN 0	-24.34	-25.03	50.44	15.00
160 LEU N		-26.69	51.52	15.00
160 LEU CA	-23.86	-26.27	52.80	15.00
160 LEU CB	-23.99	-27.45	53.77	15.00
160 LEU CG	-24.87	-28.62	53.30	15.00
160 LEU CD1	-25.08	-29.56	54.47	15.00
160 LEU CD2	-26.21	-28.16	52.76	15.00
160 LEU C	-22.91	-25.21	53.33	15.00
160 LEU O	-21.77	-25.51	53.65	15.00
161 ASN N	-23.38	-23.97	53.35	15.00
161 ASN CA	-22.55	-22.84	53.78	15.00
161 ASN CB	-22.32	-21.90	52.60	15.00
161 ASN CG	-23.61	-21.59	51.84	15.00
161 ASN OD1	-24.49	-20.86	52.31	15.00
161 ASN ND2	-23.75	-22.18	50.66	15.00
161 ASN C	-23.10	-22.04	54.94	15.00
161 ASN 0	-22.37	-21.29	55.58	15.00
162 HIS N	-24.38	-22.20	55.24	15.00
162 HIS CA	-25.01	-21.44	56.31	15.00
162 HIS CB	-26.00	-20.43	55.72	15.00
162 HIS CG		-19.38	56.68	15.00
162 HIS CD2		-18.91	56.97	15.00
162 HIS ND1	-25.58	-18.66	57.46	15.00
162 HIS CE1	-26.25	-17.79	58.19	15.00
162 HIS NE2 162 HIS C	-27.54	-17.92	57.90	15.00
162 HIS C		-22.34	57.31	15.00
		-23.07	56.94	15.00
163 ALA N		-22.30	58.57	15.00
163 ALA CA	-25.93	-23.09	59.63	15.00
163 ALA CB	~24.98	-23.25	60.31	15.00

163	ALA	С	-27.18	-22.33	60.08	15.00
163	ALA	0	-27.20	-21.09	60.08	15.00
164	VAL	N	-28.21	-23.05	60.48	15.00
164	VAL	CA	-29.44	-22.43	60.91	15.00
164	VAL	CB	-30.31	-22.10	59.66	15.00
164	VAL	CG1	-31.25	-23.23	59.31	15.00
164	VAL	CG2	-30.99	-20.79	59.85	15.00
164	VAL	С	-30.13	-23.35	61.91	15.00
164	VAL	0	-29.51	-24.32	62.35	15.00
165	LEU	N	-31.38	-23.10	62.28	15.00
165	LEU	CA	-32.05	-23.93	63.28	15.00
165	LEU	CB	-31.95	-23.25	64.64	15.00
165	LEU	CG	-32.41	-23.97	65.90	15.00
165	LEU	CD1	-31.39	-25.01	66.31	15.00
165	LEU	CD2	-32.58	-22.96	67.00	15.00
165	LEU	С	-33.51	-24.20	62.98	15.00
165	LEU	0	-34.30	-23.27	62.86	15.00
166	ALA	N	-33.88	-25.48	62.93	15.00
166	ALA	CA	-35.27	-25.89	62.66	15.00
166	ALA	CB	-35.30	-27.31	62.15	15.00
166	ALA	C	-36.07	-25.78	63.95	15.00
166	ALA	0	-35.74	-26.40	64.95	15.00
167	VAL	N	-37.14	-24.99	63.93	15.00
167	VAL	CA	-38.00	-24.75	65.09	15.00
167	VAL	CB	-38.22	-23.21	65.30	15.00
167	VAL		-39.07	-22.93	66.50	15.00
167	VAL	CG2	-36.88	-22.50	65.47	15.00
167	VAL	С	-39.35	-25.46	64.93	15.00
167	VAL	0	-40.25	-25.34	65.77	15.00
168	GLY	N	-39.51	-26.21	63.85	15.00
168	GLY	CA	-40.75	-26.91	63.61	15.00
168	GLY		-40.97	-27.18	62.14	15.00
168	GLY	0	-40.03	-27.15	61.35	15.00
	TYR		-42.22	-27.45	61.78	15.00
	TYR		-42.63	-27.75	60.41	15.00
	TYR		-42.23	-29.18	60.00	15.00
	TYR			-30.27	60.93	15.00
	TYR			-30.66	60.94	15.00
	TYR			-31.56	61.81	15.00
	TYR			-30.91	61.81	15.00
	TYR			-31.51	62.68	15.00
169			-43.63		62.68	15.00
169	TYR	CH	-44.97	-33.24	63.57	15.00

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169 TYR C	-44.14		60.31	15.00
169 TYR O	-44.81		61.30	15.00
170 GLY N	-44.70	_	59.12	15.00
170 GLY CA	-46.14	-27.63	58.97	15.00
170 GLY C	-46.56		57.52	15.00
170 GLY O	-45.85	-27.94	56.61	15.00
171 ILE N	-47.74	-26.97	57.30	15.00
171 ILE CA	-48.28	-26.78	55.97	15.00
171 ILE CB	-49.43	-27.78	55.69	15.00
171 ILE CG2	-50.14	-27.44	54.38	15.00
171 ILE CG1	-48.91	-29.22	55.69	15.00
171 ILE CD1	-48.38	-29.69	54.35	15.00
171 ILE C	-48.89	-25.40	56.05	15.00
171 ILE O	-49.85	-25.20	56.78	15.00
172 GLN N	-48.29	-24.41	55.40	15.00
172 GLN CA	-48.86	-23.07	55.47	15.00
172 GLN CB	-47.96	-22.05	54.75	15.00
172 GLN CG	-48.37	-20.59	54.97	15.00
172 GLN CD	-47.48	-19.62	54.20	15.00
172 GLN OE1	-46.94	-19.96	53.14	15.00
172 GLN NE2	<b>-47.33</b>	-18.41	54.72	15.00
172 GLN C	-50.22	-23.15	54.78	15.00
172 GLN 0	-51.26	-22.82	55.37	15.00
173 LYS N	-50.21	-23.61	53.54	15.00
173 LYS CA	-51.45	-23.77	52.79	15.00
173 LYS CB	-52.00	-22.41	52.33	15.00
173 LYS CG	-53.53	-22.35	52.25	15.00
173 LYS CD	-54.00	-20.89	52.22	15.00
173 LYS CE	-55.51	-20.75	52.34	15.00
173 LYS NZ	-55.91	-19.31	52.33	15.00
173 LYS C	-51.07	-24.63	51.61	15.00
173 LYS O	-50.50	-24.16	50.63	15.00
174 GLY N	-51.28	-25.93	51.80	15.00
174 GLY CA	-50.97	-26.89	50.76	15.00
174 GLY C	-49.51	-27.27	50.74	15.00
174 GLY O	-49.17	-28.42	50.45	15.00
175 ASN N	-48.63	-26.34	51.08	15.00
175 ASN CA	-47.20	-26.64	51.05	15.00
L75 ASN CB	-46.44	-25.52	50.34	15.00
L75 ASN CG	-46.88	-25.35	48.88	15.00
175 ASN OD1	-47.59	-24.39	48.54	15.00
175 ASN ND2	-46.49	-26.29	48.03	15.00
175 ASN C	-46.52	-26.98	52.38	15.00

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175 ASN O	-46.58	-26.21	53.35	15.00
176 LYS N	-45.88	-28.15	52.40	15.00
176 LYS CA	-45.15	-28.65	53.56	15.00
176 LYS CB	-44.68	-30.09	53.34	15.00
176 LYS CG	-45.74	-31.03	52.77	15.00
176 LYS CD	-45.35	-32.49	52.97	15.00
176 LYS CE	-46.23	-33.44	52.14	15.00
176 LYS NZ	-45.80	-33.48	50.70	15.00
176 LYS C	-43.94	-27.75	53.75	15.00
176 LYS O	-43.26	-27.41	52.78	15.00
177 HIS N	-43.64	-27.39	55.00	15.00
177 HIS CA	-42.51	-26.50	55.25	15.00
177 HIS CB	-42.96	-25.05	55.09	15.00
177 HIS CG	-43.85	-24.56	56.20	15.00
177 HIS CD2	-43.59	-24.27	57.49	15.00
177 HIS ND1	-45.16		55.99	15.00
177 HIS CE1	-45.67	-23.71	57.10	15.00
177 HIS NE2	-44.74	-23.75	58.03	15.00
177 HIS C	-41.82	-26.66	56.59	15.00
177 HIS O	-42.44	-27.05	57.57	15.00
178 TRP N	-40.52		56.60	15.00
178 TRP CA	-39.67	-26.38	57 <b>.7</b> 7	15.00
178 TRP CB	-38.26	-26.80	57.38	15.00
178 TRP CG	-38.11	-28.21	56.93	15.00
178 TRP CD2	-38.28	-29.38	57.73	15.00
178 TRP CE2	-38.02	-30.49	56.90	15.00
178 TRP CE3	-38.62	-29.60	59.08	15.00
178 TRP CD1	-37.77	-28.64	55.68	15.00
178 TRP NE1	-37.71	-30.01	55.65	15.00
178 TRP CZ2	-38.09	-31.80	57.37	15.00
178 TRP CZ3	-38.69			15.00
178 TRP CH2	-38.43	-31.98	58.68	15.00
178 TRP C	-39.59	-24.94	58.28	15.00
178 TRP O		-24.03	57.51	15.00
179 ILE N	-39.92		59.55	15.00
179 ILE CA	-39.85		60.09	15.00
179 ILE CB	-40.86	-23.17	61.23	15.00
179 ILE CG2	-40.80	-21.75	61.74	15.00
179 ILE CG1	-42.27	-23.50	60.72	15.00
179 ILE CD1	-43.38		61.70	15.00
179 ILE C		-23.13	60.55	15.00
179 ILE O		-23.79	61.47	15.00
180 ILE N	-37.73	-22.22	59.88	15.00

180 ILE CA	-36.32		60.17	15.00
180 ILE CB	-35.50		58.86	15.00
180 ILE CG2	-34.06	-21.71	59.13	15.00
180 ILE CG1	-35.63	-23.40	58.22	15.00
180 ILE CD1	-34.94	_ <del>-</del>	58.99	15.00
180 ILE C	-36.01	-20.62	60.88	15.00
180 ILE O	-36.59	-19.58	60.56	15.00
181 LYS N	-35.09	-20.68	61.84	15.00
181 LYS CA	-34.67	-19.51	62.60	15.00
181 LYS CB	-34.58	-19.83	64.09	15.00
181 LYS CG	-34.19	-18.62	64.93	15.00
181 LYS CD	-33.86	-18.95	66.37	15.00
181 LYS CE	-33.68	-17.66	67.16	15.00
181 LYS NZ	-33.07	-17.85	68.50	15.00
181 LYS C	-33.30	-19.10	62.12	15.00
181 LYS O	-32.31	-19.74	62.48	15.00
182 ASN N	-33.22	-18.04	61.32	15.00
182 ASN CA	-31.93	-17.56	60.79	15.00
182 ASN CB	-32.17	-16.80	59.48	15.00
182 ASN CG	-30.98	-16.86	58.53	15.00
182 ASN OD1	-29.87	-17.20	58.92	15.00
182 ASN ND2	-31.21	-16.50	57.27	15.00
182 ASN C	-31.29	-16.65	61.84	15.00
182 ASN O	-31.86	-16.44	62.91	15.00
183 SER N	-30.11	-16.11	61.56	15.00
183 SER CA	-29.44	-15.21	62.52	15.00
183 SER CB	-28.26	-15.93	63.18	15.00
183 SER OG	-27.41	-16.54	62.21	15.00
183 SER C	-28.98	-13.90	61.87	15.00
183 SER O	-27.84	-13.44	62.08	15.00
184 TRP N	-29.85	-13.26	61.11	15.00
184 TRP CA	-29.52	-12.01	60.45	15.00
184 TRP CB	-29.70	-12.14	58.94	15.00
184 TRP CG	-28.74	-13.05	58.26	15.00
184 TRP CD2	-28.89	-13.61	56.96	15.00
184 TRP CE2	-27.72	-14.38	56.71	15.00
184 TRP CE3	-29.87	-13.53	55.98	15.00
184 TRP CD1	-27.54	-13.48	58.74	15.00
184 TRP NE1	-26.92	-14.29	57.81	15.00
184 TRP CZ2	-27.53	-15.07	55.51	15.00
1.84 TRP CZ3	-29.68	-14.22	54.78	15.00
184 TRP CH2	-28.52	-14.98	54.56	15.00
184 TRP C	-30.45	-10.91	60.98	15.00

184	TRP	0	-31.16	-10.28	60.20	15.00
185	GLY	N	-30.48	-10.71	62.29	15.00
185	GLY	CA	-31.34	-9.69	62.86	15.00
185	GLY	С	-32.79	-10.02	62.61	15.00
185	GLY	0	-33.09	-11.01	61.96	15.00
186	GLU	N	-33.73	-9.25	63.14	15.00
186	GLU	CA	-35.13	-9.57	62.89	15.00
186	GLU	CB	-36.01	-9.41	64.14	15.00
186	GLU	CG	-36.02	-8.04	64.75	15.00
186	GLU	CD	-37.04	-7.92	65.87	15.00
186	GLU	OE1	-36.66	-7.55	67.00	15.00
186	GLU	OE2	-38.23	-8.21	65.61	15.00
186	GLU	С	-35.67	-8.78	61.72	15.00
186	GLU	0	-36.84	-8.91	61.34	15.00
187	ASN	N	-34.80	-7.96	61.13	15.00
187	ASN	CA	-35.17	-7.15	59.99	15.00
187	ASN	CB	-34.20	-5.97	59.84	15.00
187	asn	CG	-34.83	-4.76	59.13	15.00
187	ASN	OD1	-34.15	-4.03	58.42	15.00
187	asn	ND2	-36.12	-4.54	59.36	15.00
187	ASN	C	-35.15	-8.03	58.74	15.00
187	ASN	0	-35.75	-7.72	57.72	15.00
188	TRP	N	-34.45	-9.16	58.81	15.00
188	TRP	CA	-34.37	-10.07	57.66	15.00
188	TRP	CB	-33.13	-10.97	57. <i>77</i>	15.00
188	TRP	CG	-33.03	-11.92	56.63	15.00
188	TRP	CD2	-33.46	-13.29	56.62	15.00
188	TRP	CE2	-33.26	-13.77	55.31	15.00
188	TRP	CE3	-34.00	-14.15	57.58	15.00
188	TRP	CD1	-32.60	-11.64	55.38	15.00
188	TRP	NE1	-32.73	-12.74	54.58	15.00
188	TRP		-33.58	-15.08	54.93	15.00
188	TRP		-34.31	-15.45	57.20	15.00
	TRP			-15.90	55.89	15.00
	TRP			-10.94	57.55	15.00
	TRP		-36.34	-11.13	58.5 <b>4</b>	15.00
189			-35.89	-11.44	56.34	15.00
189	GLY		-37.04	-12.29	56.09	15.00
	GLY		-38.26	-11.99	56.92	15.00
	GLY		-38.56	-10.84	57.23	15.00
	ASN		-38.99	-13.02	57.32	15.00
	ASN			-12.81	58.14	15.00
190	ASN	CB	-41.17	-13.96	57.97	15.00

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190			-42.58	-13.57	58.36	15.00
190		OD1	-43.53	-13.95	57.69	15.00
190		ND2	-42.72	-12.80	59.43	15.00
190	ASN	С	-39.79	-12.68	59.61	15.00
190	ASN	0	-39.86	-13.65	60.36	15.00
191	LYS	N	-39.41	-11.47	60.02	15.00
191	LYS	CA	-39.01	-11.18	61.41	15.00
191	LYS	CB	-40.23	-11.26	62.34	15.00
191	LYS	CG	-41.41	-10.38	61.96	15.00
191	LYS	CD	-42.64	-10.75	62.79	15.00
191	LYS	CE	-43.93	-10.18	62.20	15.00
191	LYS	NZ	-45.15	-10.69	62.91	15.00
191	LYS	С	-37.94	-12.14	61.90	15.00
191	LYS	0	-38.00	-12.62	63.03	15.00
192	GLY	N	-36.97	-12.45	61.05	15.00
192	GLY	CA	-35.91	-13.35	61.43	15.00
192	GLY	С	-36.06	-14.79	60.97	15.00
192	GLY	0	-35.07	-15.51	60.83	15.00
193 '	TYR	N	-37.30	-15.21	60.70	15.00
	TYR	CA	-37.56	-16.59	60.27	15.00
	TYR	CB	-38.79	-17.14	61.01	15.00
193 !	TYR	CG	-38.57	-17.28	62.49	15.00
193	TYR	CD1	-38.77	-16.20	63.35	15.00
	ΓΥR	CE1	-38.49	-16.30	64.70	15.00
193	TYR	CD2	-38.09	-18.47	63.04	15.00
	IYR	CE2	-37.81	-18.58	64.40	15.00
	IYR	CZ	-38.00	-17.48	65.22	15.00
	TYR	OH	-37.69	-17.57	66.55	15.00
	<b>TYR</b>	C	-37.77	-16.76	58.76	15.00
	IYR	0	-37.85	-15.80	57.99	15.00
	ILE		-37.83	-18.01	58.33	15.00
	ILE		-38.06	-18.34	56.94	15.00
	ILE		-36.77	-18.17	56.07	15.00
	ILE	CG2	-35.64	-19.02	56.60	15.00
	II.E		-37.07	-18.54	54.62	15.00
	ILE	CD1	-35.87	-18.54	53.71	15.00
	[LE	C	-38.59	-19.77	56.84	15.00
	[LE	0	-38.12	-20.68	57.53	15.00
195 I	LEU :	N	-39.65	-19.93	56.06	15.00
195 I	LEU	CA	-40.27	-21.23	55.84	15.00
1.95 I	EU	CB	-41.76	-21.08	55.61	15.00
195 I	EU (	CG	-42.62	-21.03	56.88	15.00
195 I	EU (	CD1	-41.90	-20.32	58.01	15.00

195 LEU CD2	-43.94	-20.37	56. <b>5</b> 8	15.00
195 LEU C	-39.61	-21.79	54.60	15.00
195 LEU O	-39.53	-21.13	53.57	15.00
196 MET N	-39.08	-23.00	54.71	15.00
196 MET CA	-38.42	-23.63	53.58	15.00
196 MET CB	-36.96	-23.90	53.93	15.00
196 MET CG	-36.13	-22.63	54.11	15.00
196 MET SD	-34.43	-22.96	54.57	15.00
196 MET CE	-33.85	-23.77	53.09	15.00
196 MET C	-39.17	-24.90	53.20	15.00
196 MET O	-39.74	-25.58	54.05	15.00
197 ALA N	-39.23	-25.22	51.91	15.00
197 ALA CA	-39.93	-26.41	51.45	15.00
197 ALA CB	-39.79	-26.55	49.94	15.00
197 ALA C	-39.51	-27.70	52.15	15.00
197 ALA O	-38.32	-27.92	52.41	15.00
198 ARG N	-40.50	-28.53	52.47	15.00
198 ARG CA	-40.29	-29.81	53.12	15.00
198 ARG CB	-40.95	-29.84	54.50	15.00
198 ARG CG	-40.91	-31.22	55.15	15.00
198 ARG CD	-41.22	_	56.63	15.00
198 ARG NE	-42.59		56.98	15.00
198 ARG CZ	-43.60	-31.70	56.98	15.00
198 ARG NH1	-43.40	-32.96	56.63	15.00
198 ARG NH2	-44.79	-31.32	57.43	15.00
198 ARG C	-40.86		52.27	15.00
198 ARG O	-42.03	-30.91	51.88	15.00
199 ASN N	-40.00	-31.90	51.96	15.00
199 ASN CA	-40.33	-33.08	51.17	15.00
199 ASN CB	-41.68	-33.69	51.58	15.00
199 ASN CG	-41.66			15.00
199 ASN OD1	-42.63		53.74	15.00
199 ASN ND2	-40.55	-34.91	53.38	15.00
199 ASN C		-32.89	49.66	15.00
199 ASN O		-33.83	48.89	15.00
200 LYS N		-31.71	49.23	15.00
200 LYS CA	-39.60	-31.45	47.81	15.00
200 LYS CB	-39.83	-29.98	47.45	15.00
200 LYS CG	-41.27	-29.52	47.67	15.00
200 LYS CD		-28.27	46.88	15.00
200 LYS CE	-43.08	-27.86	47.07	15.00
200 LYS NZ		-26.73	46.16	15.00
200 LYS C	-38.15	-31.86	47.56	15.00

200 LYS O	-37.29		47.31	15.00
201 ASN N	-37.90		47.74	15.00
201 ASN CA	-36.58	-33.74	47.55	15.00
201 ASN CB	-36.23	-33.86	46.06	15.00
201 ASN CG	-37.27	-34.62	45.25	15.00
201 ASN OD1	-37.46	-34.34	44.06	15.00
201 ASN ND2	-37.96	-35.58	45.88	15.00
201 ASN C	-35.48	-32.95		15.00
201 ASN O	-34.88	-32.07	47.64	15.00
202 ASN N	-35.27	-33.23	49.53	15.00
202 ASN CA	-34.21	-32.62	50.34	15.00
202 ASN CB	-33.01	-33.56	50.33	15.00
202 ASN CG	-31.96	-33.20	51.34	15.00
202 ASN OD1	-32.27	-32.76	52.45	15.00
202 ASN ND2	-30.71		50.98	15.00
202 ASN C	-33.81	-31.18	49.94	15.00
202 ASN 0	-32.63	-30.88	49.70	15.00
203 ALA N	-34.78		49.91	15.00
203 ALA H	-35.58	-30.58	50.39	15.00
203 ALA CA	-34.60		49.49	15.00
203 ALA CB	-35.83	-28.06	49.83	15.00
203 ALA C	-33.41	-28.27	50.25	15.00
203 ALA O	-33.31	-28.25	51.47	15.00
204 CYS N	-32.47	-27.71	49.46	15.00
204 CYS CA	-31.31	-27.02	50.02	15.00
204 CYS C	-30.43	-27.87	50.92	15.00
204 CYS 0	-29.53	-27.34	51.59	15.00
204 CYS CB	-31.78	-25.79	50.79	15.00
204 CYS SG	-32.67		49.75	15.00
205 GLY N	-30.64	-29.19	50.92	15.00
205 GLY CA	-29.86	-30.06	51.76	15.00
205 GLY C	-30.07	-29.75	53.23	15.00
205 GLY O	-29.14	-29.81	54.04	15.00
206 ILE N	-31.29	-29.37	53.59	15.00
206 ILE CA	-31.60	-29.06	54.98	15.00
206 ILE CB	-33.08	-28.64	55.16	15.00
206 ILE CG2	-34.01	-29.71	54.60	15.00
206 ILE CG1	-33.38	-28.40	56.63	15.00
206 ILE CD1	-34.65	-27.65	56.89	15.00
206 ILE C	-31.30	-30.24	55.91	15.00
206 ILE 0		-30.05	57.07	15.00
207 ALA N	-31.42	-31.46	55.39	15.00
207 ALA CA	-31.16	-32.64	56.19	15.00

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207	ALA	CB	-32.30	-33.64	56.03	15.00
207	ALA	С	-29.82	-33.31	55.92	15.00
207	ALA	. 0	-29.71	-34.52	55.99	15.00
208	asn	N	-28.75	-32.54	55.69	15.00
208	asn	CA	-27.44	-33.14	55.41	15.00
208	ASN	CB	-26.91	-32.66	54.07	15.00
208	ASN	CG	-27.50	-33.42	52.92	15.00
208	ASN	OD1	-28.69	-33.32	52.66	15.00
208	ASN	ND2	-26.68	-34.22	52.24	15.00
208	ASN	С	-26.38	-32.91	56.46	15.00
208	ASN	0	-25.30	-33.49	56.40	15.00
209	LEU	N	-26.65	-32.01	57.40	15.00
209	LEU	CA	-25.71	-31.72	58.47	15.00
209	LEU	CB	-24.78	-30.58	58.07	15.00
209	LEU	CG	-23.44	-30.57	58.80	15.00
209	LEU	CD1	-22.65	-31.81	58.41	15.00
209	LEU	CD2	-22.65	-29.33	58.46	15.00
209	LEU	С	-26.52	-31.34	59.69	15.00
209	LEU	0	-26.24	-30.34	60.34	15.00
210	ALA	N	-27.55	-32.13	59.94	15.00
210	ALA	H	-27.87	-32.61	59.15	15.00
210	ALA	CA	-28.41	-31.91	61.10	15.00
210	ALA	CB	-29.82	-32.45	60.85	15.00
210	ALA	С	-27.84	-32.66	62.31	15.00
210	ALA	0	-27.12	-33.64	62.12	15.00
211	SER	N	-28.10	-32.14	63.49	15.00
211	SER	CA	-27.62	-32.76	64.72	15.00
211	SER	CB	-26.11	-32.53	64.90	15.00
211	SER	OG	-25.80	-31.16	65.02	15.00
211	SER	С	-28.35	-32.12	65.88	15.00
211	SER	0	-28.97	-31.08	65.73	15.00
212	PHE	N	-28.31	-32.77	67.03	15.00
212	PHE	CA	-28.95	-32.21	68.21	15.00
212	PHE	CB	-30.43	-32.63	68.32	15.00
212	PHE	CG	-30.66	-34.13	68.43	15.00
212	PHE	CD1	-31.05	-34.86	67.32	15.00
212	PHE	CD2	-30.55	-34.79	69.67	15.00
212	PHE	CE1	-31.34	-36.23	67.44	15.00
212	PHE	CE2	-30.83	-36.15	69.79	15.00
212	PHE	CZ	-31.23	-36.86	68.67	15.00
212	PHE	С		-32.62	69.42	15.00
212	PHE	0		-33.67	69.42	15.00
213	PRO	N ·		-31.73	70.40	15.00

		IABLE	ш	
213 PRO CD	-28.49	-30.34	70.46	15.00
213 PRO CA	-27.25	-32.08	71.59	15.00
213 PRO CB	-26.99	-30.73	72.23	15.00
213 PRO CG	-28.24		71.91	15.00
213 PRO C	-28.08	-32.96	72.51	15.00
213 PRO 0	-29.31	-32.92		15.00
214 LYS N	-27.43	-33.79	73.30	15.00
214 LYS CA	-28.14	-34.63	74.23	15.00
214 LYS CB	-27.49	-36.02	74.28	15.00
214 LYS CG	-28.32	-37.14	73.61	15.00
214 LYS CD	-27.43	-38.10	72.81	15.00
214 LYS CE	-26.30	-38.66	73.65	15.00
214 LYS NZ	-25.35	-39.45	72.81	15.00
214 LYS C	-28.02	-33.93	75.57	15.00
214 LYS O	-26.95	-33.45	75.91	15.00
215 MET N	-29.11	-33.79	76.30	15.00
215 MET CA	-29.01	~33.15	77.60	15.00
215 MET CB	-29.60	-31.73	77.56	15.00
215 MET CG	-28.77	-30.73	78.36	15.00
215 MET SD	-29.41	-29.06	78.37	15.00
215 MET CE	-30.41	-29.07	79.82	15.00
215 MET C	-29.66	-33.98	78.71	15.00
215 MET OT1	-30.59	-34.77	78.41	15.00
215 MET OT2	-29.20	-33.85	79.87	15.00
216 нон он2	-28.05	-18.06	84.86	15.00
217 НОН ОН2	-23.19	-33.36	81.36	15.00
218 НОН ОН2	-31.64	-15.80	65.41	15.00
219 НОН ОН2	-30.17	-19.91	64.18	15.00
220 нон он2	-13.36	-11.60	62.86	15.00
221 нон он2	-9.95	-9.46	71.42	15.00
222 НОН ОН2	-34.59	-22.68	70.30	15.00
223 НОН ОН2	-17.52	-33.99	64.33	15.00
224 НОН ОН2	-15.72	-11.02	61.35	15.00
225 НОН ОН2	-24.41	-30.51	62.51	15.00
226 нон он2	-10.27	-5.38	68.19	15.00
227 НОН ОН2		-16.84	67.70	15.00
228 НОН ОН2		-30.73	49.92	15.00
229 нон он2		-36.65	56.24	15.00
230 нон он2	-37.78	-15.40	68.33	15.00
231 нон он2	-38.40	-35.66	51.51	15.00
232 нон он2		-36.36	61.93	15.00
233 нон он2	-41.75	-34.32	46.57	15.00
234 НОН ОН2	-28.01	-19.38	62.11	15.00

235 НОН ОН2	-21 94	-29.60	60 55	15 00
236 НОН ОН2	-26.15			
237 НОН ОН2	-29.39			15.00
238 HOH OH2	-30.20	-20.71	79.14	15.00
239 НОН ОН2	-40.59	-13.37	84.30	15.00
240 HOH OH2	-36.04		49.72	15.00
241 HOH OH2	-46.35			15.00
242 HOH OH2	-24.71		56.72 61.99	15.00
243 HOH OH2	-44.08		65.62	15.00
244 HOH OH2	-25.57		65.10	15.00
245 HOH OH2	-33.44		71.31	15.00
246 HOH OH2	-47.48		77.05	15.00
247 HOH OH2	-14.60	-14.01	81.32	15.00
248 HOH OH2	-7.93	-18.05	73.48	15.00
249 HOH OH2	-7.49	-16.70	75.98	15.00 15.00
250 НОН ОН2	-26.27		59.26	15.00
251 HOH OH2	-35.15		53.73	15.00
252 НОН ОН2	-33.62	-27.20	46.10	15.00
253 нон он2	-40.60	-16.73	45.07	15.00
254 HOH OH2	-41.25	-34.55	55.94	15.00
255 нон он2	-40.71		72.64	15.00
256 нон он2	-32.67	-13.41	60.76	15.00
257 НОН ОН2	-39.61	-9.04	58.76	15.00
258 нон он2	-31.33	-8.54	65.90	15.00
259 НОН ОН2	-31.41	-5.90	63.60	15.00
260 HOH OH2	-19.54	-8.02	63.36	15.00
261 HOH OH2	-33.59	-19.88	70.38	15.00
262 НОН ОН2	-32.78	-42.12	66.81	15.00
263 нон он2	-13.22	-22.75	77.99	15.00
264 HOH OH2	-8.15	-22.46	73.27	15.00
265 нон он2	-9.06	-29.93	75.92	15.00
266 НОН ОН2	-20.77	-33.56	62.36	15.00
267 НОН ОН2	-24.27	-45.12	64.98	15.00
268 НОН ОН2	-11.63		70.13	15.00
269 HOH OH2	-11.87	-26.78	72.10	15.00
270 нон он2		-34.47	79.41	15.00
271 нон он2		-37.69	70.75	15.00
272 НОН ОН2		-24.81	88.02	15.00
273 НОН ОН2		-12.16	66.61	15.00
274 HOH OH2			69.94	15.00
275 HOH OH2		-20.23	55.91	15.00
276 НОН ОН2		-20.58	48.90	15.00
277 НОН ОН2	-25.90	-22.48	45.16	15.00

WO 97/16177		PCT/US96/17512
	TABLE III	
278 HOH OH2 279 HOH OH2 280 HOH OH2 281 HOH OH2	-20.12 -27.71 -38.87 -31.38	52.91 15.00 33.74 15.00 31.99 15.00 31.07 15.00

6	ASP	CA	-43.81	-24.70	66.29	15.00
6	ASP	CB	-44.76	-23.60	65.80	15.00
6	ASP	CG	-44.06	-22.25	65.59	15.00
6	ASP	OD1	-42.88	-22.10	65.94	15.00
6	ASP	OD2	-44.73	-21.33	65.08	15.00
6	ASP	С	-43.41	-24.48	67.75	15.00
6	ASP	0	-44.26	-24.33	68.63	15.00
7	TYR	N	-42.12	-24.54	68.00	15.00
7	TYR	CA	-41.60	-24.37	69.34	15.00
7	TYR	CB	-40.20	-24.96	69.42	15.00
7	TYR	CG	-40.23	-26.49	69.41	15.00
7	TYR	CD1	-40.62	-27.20	70.55	15.00
7	TYR	CE1	-40.66	-28.57	70.55	15.00
7	TYR	CD2	-39.89	-27.21	68.27	15.00
7	TYR	CE2	-39.94	-28.60	68.26	15.00
7	TYR	CZ	-40.32	-29.27	69.41	15.00
7	TYR	OH	-40.40	-30.63	69.42	15.00
7	TYR	С	-41.64	-22.94	69.83	15.00
7	TYR	0	-41.52	-22.68	71.03	15.00
8	ARG	N	-41.85	-22.01	68.90	15.00
8	ARG	CA	-41.91	-20.58	69.22	15.00
8			-42.07	-19.74	67.95	15.00
8	ARG	CG	-40.84	-19.78	67.04	15.00
8	ARG	CD	-41.01	-18.96	65.78	15.00
8	ARG	NE	-41.97	-19.57	64.86	15.00
8	ARG		-42.43	-18.97	63.77	15.00
8	ARG	NH1	-42.03	-17.75	63.46	15.00
8	ARG	NH2	-43.30	-19.60	62.98	15.00
8	ARG		-43.09	-20.36	70.15	15.00
8	ARG		-42.95	-19.71	71.18	15.00
9	LYS		-44.23	-20.95	69.82	15.00
9	LYS		-45.41	-20.80	70.64	15.00
9	LYS		-46.59	-21.48	69.96	15.00
	LYS			-20.90	68.60	15.00
	LYS			-21.89	67.79	15.00
9			-48.07	-21.38	66.41	15.00
9	LYS	NZ	-48.46	-22.50	65.49	15.00
9		С	-45.16	-21.46	71.99	15.00
9	LYS	С	-45.70	-21.01	73.01	15.00
10		N	-44.29	-22.46	71.99	15.00
10				-23.25	73.18	15.00
	LYS			-24.60	72.77	15.00
10	LYS	CG	-44.19	-25.43	71.81	15.00

		IABLE	V1	
10 LYS CD	-45.03	-26.48	72.52	15.00
10 LYS CE	-46.23	-25.86	73.25	15.00
10 LYS NZ	-47.28	-25.31	72.33	15.00
10 LYS C	-42.97	-22.59	74.15	15.00
10 LYS O	-42.91	-22.97	75.32	15.00
11 GLY N	-42.15	-21.66	73.67	15.00
11 GLY CA	-41.20	-21.02	74.57	15.00
11 GLY C	-39.83	-21.68	74.55	15.00
11 GLY O	-39.00	-21.42	75.42	15.00
12 TYR N	-39.57	-22.49	73.53	15.00
12 TYR CA	-38.29	-23.18	73.39	15.00
12 TYR CB	-38.48	-24.53	72.68	15.00
12 TYR CG	-39.09	-25.67	73.49	15.00
12 TYR CD1	-40.24	-25.49	74.27	15.00
12 TYR CE1	-40.82	-26.55	74.95	15.00
12 TYR CD2	-38.55	-26.95	73.42	15.00
12 TYR CE2	-39.13	-28.01	74.09	15.00
12 TYR CZ	-40.26	-27.81	74.85	15.00
12 TYR OH	-40.86	~28.88	75.47	15.00
12 TYR C	-37.31	-22.37	72.55	15.00
12 TYR O	-36.15	-22.73	72.44	15.00
13 VAL N	-37.78	-21.29	71.94	15.00
13 VAL CA	-36.94	-20.46	71.07	15.00
13 VAL CB	-37.56	-20.37	69.65	15.00
13 VAL CG1	-36.60	-19.70	68.68	15.00
13 VAL CG2	-37.91	-21.76	69.16	15.00
13 VAL C	-36.75	-19.06	71.62	15.00
13 VAL O	-37.70	-18.41	72.02	15.00
14 THR N	-35.51	-18.61	71.66	15.00
14 THR CA	-35.21	-17.27	72.15	15.00
14 THR CB	-33.80	-17.21	72.74	15.00
14 THR OG1	-32.85	-17.58	71.74	15.00
14 THR CG2	-33.69	-18.13	73.92	15.00
14 THR C	-35.31	-16.23	71.02	15.00
14 THR O	-35.46	-16.59	69.85	15.00
15 PRO N	-35.25	-14.94	71.35	15.00
15 PRO CD	-35.15	-14.35	72.71	15.00
15 PRO CA	-35.34	-13.89	70.34	15.00
15 PRO CB	-35.15	-12.62	71.16	15.00
15 PRO CG	-35.72	-12.99	72.50	15.00
15 PRO C	-34.26	-14.00	59.25	15.00
15 PRO O	-33.13	-14.41	69.53	15.00
16 VAL N	-34.61	-13.62	68.02	15.00

16	VAL	CA	-33.69	-13.67	66.89	15.00
16	VAL	CB	-34.39	-13.43	65.54	15.00
16	VAL	CG1	-33.36	-13.27	64.43	15.00
16	VAL	CG2	-35.29	-14.58	65.20	15.00
16	VAL	C	-32.56	-12.66	67.05	15.00
16	VAL	0	-32.79	-11.47	67.28	15.00
17	LYS	N	-31.34	-13.17	66.92	15.00
17	LYS	CA	-30.15	-12.36	67.04	15.00
17	LYS	CB	-29.13	-13.09	67.91	15.00
17	LYS	CG	-29.67	-13.49	69.27	15.00
17	LYS	CD	-30.28	-12.30	69.96	15.00
17	LYS	CE	-30.93	-12.65	71.28	15.00
17	LYS	NZ	-31.68	-11.47	71.83	15.00
17	LYS	С	-29.58	-12.08	65.65	15.00
17	LYS	0	-30.13	-12.52	64.64	15.00
18	ASN	N	-28.48	-11.33	65.60	15.00
18	ASN	CA	-27.82	-10.98	64.34	15.00
18	ASN	CB	-28.02	-9.49	64.05	15.00
18	ASN	CG	-27.42	-9.05	62.72	15.00
18	ASN	OD1	-26.35	-9.49	62.32	15.00
18	ASN	ND2	-28.11	-8.15	62.03	15.00
18	ASN	С	-26.32	-11.27	64.49	15.00
18	ASN	0	-25.67	-10.71	65.37	15.00
19	GLN	N	-25.79·	-12.14	63.63	15.00
19	GLN	CA	-24.38	-12.49	63.68	15.00
19	GLN	CB	-24.08	-13.76	62.87	15.00
19	GLN	CG	-24.55	-13.74	61.41	15.00
19	GLN	CD	-24.19	-15.02	60.65	15.00
19	GLN	OE1	-25.06	-15.85	60.37	15.00
19	GLN	NE2	-22.92	-15.16	60.28	15.00
19	GLN	С	-23.43	-11.34	63.29	15.00
19	GLN	0	-22.26	-11.34	63.67	15.00
20	GLY	N	-23.92	-10.39	62.50	15.00
20	GLY	CA	-23.11	-9.25	62.11	15.00
20	GLY	C	-22.23	-9.49	60.91	15.00
20	GLY	0	-22.71	-9.92	59.87	15.00
21	GLN	N	-20.96	-9.14	61.02	15.00
21	GLN	CA	-20.02	-9.34	59.92	15.00
21	GLN	CB	-19.11	-8.11	59.75	15.00
21	GLN	CG	-19.78	-6.87	59.13	15.00
21	GLN	CD	-20.32	-7.11	57.72	15.00
21	GLN	OE1	-19.57	-7.44	56.79	15.00
21	GLN	NE2	-21.62	-6.95	57.55	15.00

	•	IABLE	A.1	
21 GLN C	-19.18	-10.59	60.17	15.00
21 GLN 0	-18.39	-11.01	59.33	15.00
22 CYS N	-19.34	-11.17	61.36	15.00
22 CYS CA	-18.63	-12.38	61.75	15.00
22 CYS C	-19.40	-13.60	61.22	15.00
22 CYS O	-20.64	-13.58	61.15	15.00
22 CYS CB	-18.52	-12.41	63.27	15.00
22 CYS SG	-17.94	-13.95	64.05	15.00
23 GLY N	-18.68	-14.63	60.78	15.00
23 GLY CA	-19.32	-15.82	60.25	15.00
23 GLY C	-19.52	-16.87	61.32	15.00
23 GLY O	-19.06	-18.00	61.18	15.00
24 SER N	-20.24	-16.49	62.37	15.00
24 SER CA	-20.52	-17.34	63.52	15.00
24 SER CB	-20.42	-16.49	64.77	15.00
24 SER OG	-21.27	-15.36	64.65	15.00
24 SER C	-21.92	-17.95	63.44	15.00
24 SER O	-22.54	-18.24	64.47	15.00
25 CYS N	-22.40	-18.17	62.23	15.00
25 CYS CA	-23.72	-18.75	62.04	15.00
25 CYS CB	-24.08	-18.74	60.55	15.00
25 CYS SG	-23.06	-19.79	59.52	15.00
25 CYS C	-23.81	-20.15	62.66	15.00
25 CYS O	-24.90	-20.59	63.05	15.00
25 INH C1	-27.01	-9.79	58.47	15.00
25 INH C2	-26.33	-10.46	59.49	15.00
25 INH C3	-25.12	-11.10	59.22	15.00
25 INH C4	-24.57	-11.08	57.94	15.00
25 INH C5	-25.26	-10.40	56.92	15.00
25 INH C6	-26.47	-9.76	57.18	15.00
25 INH C7	-23.25	-11.75	57.65	15.00
25 INH 08	-23.16	-13.18	57.51	15.00
25 INH C9	-22.82	-13.83	56.29	15.00
25 INH C10	-22.10	-13.30	55.22	15.00
25 INH C11	-23.03	-15.93	55.08	15.00
25 INH C12	-22.32	-15.39	54.01	15.00
25 INH C13	-21.85	-14.07	54.07	15.00
25 INH C14	-23.54	-17.29	55.29	15.00
25 INH 015	-24.49	-17.82	54.70	15.00
25 INH N16	-22.71	-17.77	56.29	15.00
25 INH N17	-22.76	-19.07	56.92	15.00
25 INH C18	-23.27	-15.13	56.20	15.00
25 INH C19	-22.05	-19.01	58.26	15.00

25	INH	020	-21.78	-17.83	58.57	15.00
25	INH	C21	-21.27	-30.33	52.84	15.00
25	INH	C22	-20.95	-30.49	54.19	15.00
25	INH	C23	-20.34	-29.44	54.89	15.00
25	INH	C24	-20.03	-28.23	54.25	15.00
25	INH	C25	-20.35	-28.09	52.90	15.00
25	INH	C26	-20.96	-29.12	52.19	15.00
25	INH	C27	-19.35	-27.11	55.01	15.00
25	INH	028	-20.01	-25.85	55.20	15.00
25	INH	C29	-20.09	-25.30	56.50	15.00
25	INH	030	-19.34	-25.70	57.40	15.00
25	INH	C31	-21.28	-23.64	57.93	15.00
25	INH	C32	-21.14	-24.56	59.14	15.00
25	INH	C33	-22.16	-25.68	59.35	15.00
25	INH	C34	-23.25	-25.62	58.28	15.00
25	INH	C35	-21.45	-27.01	59.33	15.00
25	INH	C36	-20.52	-22.34	58.22	15.00
25	INH	037	-19.37	-22.35	58.66	15.00
25	INH	N38	-21.23	-21.24	57.98	15.00
25	INH	N39	-20.81	-19.86	58.17	15.00
25	INH	N40	-21.01	-24.34	56.66	15.00
26	TRP	N	-22.67	-20.83	62.82	15.00
26	TRP	CA	-22.65	-22.16	63.44	15.00
26	TRP	CB	-21.35	-22.91	63.12	15.00
26	TRP	CG	-20.11	-22.22	63.59	15.00
26	TRP	CD2	-19.48	-22.37	64.87	15.00
26	TRP	CE2	-18.42	-21.44	64.92	15.00
26	TRP	CE3	-19.71	-23.18	65.98	15.00
26	TRP	CD1	-19.41	-21.27	62.93	15.00
26	TRP	NE1	-18.40	-20.78	63.72	15.00
26	TRP	CZ2	-17.59	-21.30	66.03	15.00
26	TRP	CZ3	-18.88	-23.05	67.10	15.00
	TRP		-17.84	-22.11	67.11	15.00
26	TRP	С	-22.85	-22.06	64.96	15.00
	TRP		-23.57	-22.86	65.55	15.00
	ALA	N	-22.24	-21.04	65.57	15.00
27			-22.33	-20.83	67.01	15.00
27	ALA	CB	-21.35	-19.78	67.46	15.00
27	ALA		-23.74	-20.47	67.45	15.00
	ALA	0	-24.21	-20.91	68.50	15.00
28	PHE	N	-24.42	-19.66	66.66	15.00
28	PHE	CA		-19.27	66.96	15.00
28	PHE	CB	-26.23	-18.10	66.07	15.00

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28 PHE CG	-25.67		66.49	15.00
28 PHE CD1	-24.46	-16.32	65.99	15.00
28 PHE CD2	-26.35	-15.98	67.42	15.00
28 PHE CE1	-23.92	-15.11	66.41	15.00
28 PHE CE2	-25.81	-14.78	67.84	15.00
28 PHE CZ	-24.60	-14.35	67.34	15.00
28 PHE C	-26.74	-20.47	66.82	15.00
28 PHE O	-27.62	-20.68	67.66	15.00
29 SER N	-26.56	-21.25	65.78	15.00
29 SER CA	-27.40	-22.41	65.55	15.00
29 SER CB	-27.05	-23.08	64.23	15.00
29 SER OG	-27.68	-24.35	64.15	15.00
29 SER C	-27.28	-23.44	66.66	15.00
29 SER O	-28.27	-24.06	67.03	15.00
30 SER N	-26.06	-23.65	67.16	15.00
30 SER CA	-25.79	-24.61	68.22	15.00
30 SER CB	-24.29	-24.72	68.44	15.00
30 SER OG	-23.64	-25.04	67.22	15.00
30 SER C	-26.44	-24.15	69.51	15.00
30 SER O	-27.07	-24.93	70.25	15.00
31 VAL N	-26.25	-22.87	69.80	15.00
31 VAL CA	-26.81	-22.23	70.98	15.00
31 VAL CB	-26.39	-20.75	71.00	15.00
31 VAL CG1	-27.52	-19.85	71.44	15.00
31 VAL CG2	-25.18	-20.58	71.92	15.00
31 VAL C	-28.32	-22.41	70.92	15.00
31 VAL O	-28.95	-22.72	71.94	15.00
32 GLY N	-28.89	-22.27	69.73	15.00
32 GLY CA	-30.32	-22.44	69.56	15.00
32 GLY C	-30.76	-23.83	69.97	15.00
32 GLY O	-31.77	-24.00	70.65	15.00
33 ALA N	-30.00	-24.83	69.55	15.00
33 ALA CA	-30.28	-26.21	69.89	15.00
33 ALA CB	-29.29	-27.14	69.22	15.00
33 ALA C	-30.20	-26.34	71.42	15.00
33 ALA O	-31.19	-26.70	72.06	15.00
34 LEU N	-29.06	-25.98	71.99	15.00
34 LEU CA	-28.87	-26.06	73.44	15.00
34 LEU CB	-27.55	-25.42	73.85	15.00
34 LEU CG	-26.25	-26.10	73.41	15.00
34 LEU CD1	-25.07	-25.21	73.74	15.00
34 LEU CD2	-26.11	-27.45	74.08	15.00
34 LEU C	-30.02	-25.41	74.21	15.00

34	LEU	0	-30.59	-26.01	75.13	15.00
35	GLU	N	-30.39	-24.20	73.80	15.00
35	GLU	CA	-31.46	-23.44	74.44	15.00
35	GLU	CB	-31.63	-22.08	73.77	15.00
35	GLU	CG	-30.41	-21.18	73.87	15.00
35	GLU	CD	-30.58	-19.86	73.15	15.00
35	GLU	OE1	-31.46	-19.76	72.27	15.00
35	GLU	OE2	-29.83	-18.92	73.46	15.00
35	GLU	С	-32.79	-24.17	74.42	15.00
35	GLU	0	-33.51	-24.18	75.42	15.00
36	GLY	N	-33.11	-24.77	73.27	15.00
36	GLY	CA	-34.35	-25.52	73.13	15.00
36	GLY	С	-34.42	-26.67	74.11	15.00
36	GLY	0	-35.48	-26.98	74.65	15.00
37	GLN	N	-33.28	-27.30	74.37	15.00
37	GLN	CA	-33.21	-28.42	75.29	15.00
37	GLN	CB	-31.94	-29.22	75.05	15.00
37	GLN	CG	-32.00	-30.06	73.80	15.00
37	GLN	CD	-33.19	-31.00	73.80	15.00
37	GLN	OE1	-33.32	-31.84	74.69	15.00
37	GLN	NE2	-34.07	-30.83	72.84	15.00
37	GLN	С	-33.28	-27.96	76.74	15.00
37	GLN	0	-33.94	-28.58	77.58	15.00
38	LEU	N	-32.64	-26.83	77.01	15.00
38	LEU	CA	-32.62	-26.25	78.35	15.00
38	LEU	CB	-31.77	-24.98	78.37	15.00
38	LEU	CG	-31.54	-24.37	79.75	15.00
38	LEU	CD1	-30.73	-25.34	80.61	15.00
38	LEU	CD2	-30.82	-23.05	79.63	15.00
38	LEU	С	-34.04	-25.95	78.83	15.00
38	LEU	0	-34.31	-25.95	80.02	15.00
39	LYS	N	-34.94	-25.65	77.90	15.00
39	LYS	CA	-36.32	-25.38	78.26	15.00
39	LYS	CB	-37.04	-24.66	77.12	15.00
39	LYS	CG	-38.53	-24.45	77.32	15.00
39	LYS	CD	-38.85	-23.47	78.43	15.00
39	LYS	CE	-40.35		78.70 、	15.00
39	LYS	NZ	-40.71	-22.74	79.94	15.00
39	LYS	C	-36.98	-26.72	78.54	15.00
39	LYS	0	-37.63	-26.90	79.57	15.00
40	LYS	N	-36.73	-27.68	77.65	15.00
40	LYS	CA	-37.28	-29.03	77.72	15.00
40	LYS	CB	-36.61	-29.90	76.66	15.00

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46	LEU	N	-33.18	-19.57	78.58	15.00
46	LEU	CA	-32.16	-18.56	78.80	15.00
46	LEU	CB	-31.30	-18.93	80.01	15.00
46	LEU	CG	-30.51	-17.78	80.61	15.00
46	LEU	CD1	-31.46	-16.70	81.06	15.00
46	LEU	CD2	-29.69	-18.27	81.78	15.00
46	LEU	C	-31.27	-18.40	77.56	15.00
46	LEU	0	-31.02	-19.36	76.85	15.00
47	ASN	N	-30.83	-17.17	77.29	15.00
47	ASN	CA	-29.98	-16.89	76.13	15.00
47	ASN	СВ	-29.92	-15.38	75.84	15.00
47	ASN	CG	-31.27	-14.80	75.53	15.00
47	ASN	OD1	-31.93	-14.22	76.41	15.00
47	ASN	ND2	-31.71	-14.94	74.29	15.00
47	ASN	С	-28.57	-17.36	76.39	15.00
47	ASN	0	-27.91	-16.86	77.30	15.00
48	LEU	N	-28.10	-18.32	75.61	15.00
48	LEU	CA	-26.75	-18.84	75.78	15.00
48	LEU	CB	-26.70	-20.31	75.39	15.00
48	LEU	CG	-27.60	-21.19	76.26	15.00
48	LEU	CD1	-27.18	-22.64	76.10	15.00
48	LEU	CD2	-27.50	-20.77	77.73	15.00
48	LEU	C	-25.77	-18.04	74.98	15.00
48	LEU	0	-26.14	-17.34	74.04	15.00
49	SER	N	-24.50	-18.13	75.34	15.00
49	SER	CA	-23.47	-17.36	74.67	15.00
49	SER	CB	-22.34	-17.06	75.64	15.00
49	SER	OG	-21.34	-16.25	75.03	15.00
49	SER	C	-22.89	-17.98	73.40	15.00
49	SER	0	-22.29	-19.06	73.45	15.00
50	PRO	N	-23.07	-17.31	72.24	15.00
50	PRO	CD -	-24.05	-16.24	72.05	15.00
50	PRO	CA	-22.55	-17.77	70.95	15.00
50	PRO	CB	-23.37	-16.96	69.95	15.00
50	PRO	CG	-24.61	-16.59	70.71	15.00
50	PRO	С	-21.09	-17.37	70.86	15.00
50	PRO	0	-20.29	-18.01	70.19	15.00
51	GLN	N	-20.74	-16.27	71.52	15.00
	GLN		-19.37	-15.78	71.56	15.00
51	GLN	CB	-19.30	-1.4.45	72.33	15.00
51	GLN	CG	-17.93	-13.77	72.34	15.00
51	GLN	CD	-17.55	-13.17	71.00	15.00
51	GLN	OE1	-18.39	-12.57	70.31	15.00

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51 GLN NE2	-16.29	-13.29	70.64	15.00
51 GLN C	-18.53	-16.82	72.26	15.00
51 GLN O	-17.45	-17.17	71.80	15.00
52 ASN N	-19.03	-17.33	73.38	15.00
52 ASN CA	-18.30	-18.34	74.14	15.00
52 ASN CB	-19.20	-18.91	75.24	15.00
52 ASN CG	-18.51	-19.98	76.10	15.00
52 ASN OD1	-19.13	-20.54	76.99	15.00
52 ASN ND2	-17.23	-20.22	75.86	15.00
52 ASN C	-17.86	-19.45	73.19	15.00
52 ASN O	-16.70	-19.87	73.22	15.00
53 LEU N	-18.76	-19.88	72.32	15.00
53 LEU CA	-18.42	-20.92	71.38	15.00
53 LEU CB	-19.66	-21.39	70.63	15.00
53 LEU CG	-20.68	-22.14	71.49	15.00
53 LEU CD1	-21.63	-22.88	70.59	15.00
53 LEU CD2	-19.98	-23.13	72.38	15.00
53 LEU C	-17.35	-20.41	70.42	15.00
53 LEU O	-16.28	-21.02	70.31	15.00
54 VAL N	-17.61	-19.27	69.79	15.00
54 VAL CA	-16.68	-18.63	68.86	15.00
54 VAL CB	-17.16	-17.18	68.52	15.00
54 VAL CG1	-16.02	-16.35	67.96	15.00
54 VAL CG2	-18.31	-17.21	67.54	15.00
54 VAL C	-15.24	-18.57	69.37	15.00
54 VAL O	-14.31	-18.91	68.66	15.00
55 ASP N	-15.07	-18.14	70.61	15.00
55 ASP CA	-13.75	-18.00	71.21	15.00
55 ASP CB	-13.78	-16.96	72.33	15.00
55 ASP CG	-14.29	-15.61	71.87	15.00
55 ASP OD1	-14.16	-15.30	70.67	15.00
55 ASP OD2	-14.79	-14.85	72.72	15.00
55 ASP C	-13.10	-19.26	71.77	15.00
55 ASP 0	-11.88	-19.33	71.86	15.00
56 CYS N	-13.89	-20.25	72.17	15.00
56 CYS CA	-13.31	-21.45	72.77	15.00
56 CYS C	-13.25	-22.75	71.96	15.00
56 CYS 0	-12.44	-23.62	72.27	15.00
56 CYS CB	-13.98	-21.73	74.11	15.00
56 CYS SG	-14.30	-20.24	75.11	15.00
57 VAL N	-14.09	-22.89	70.94	15.00
57 VAL CA	-14.08	-24.13	70.16	15.00
57 VAL CB	-15.43	-24.34	69.42	15.00

57 VAL CG1       -15.47       -25.73       68.8         57 VAL CG2       -16.59       -24.17       70.3         57 VAL C       -12.91       -24.19       69.1         57 VAL O       -13.08       -24.04       67.9         58 SER N       -11.71       -24.46       69.7         58 SER CA       -10.50       -24.55       68.8         58 SER CB       -9.34       -25.09       69.7         58 SER CB       -9.08       -24.27       70.8         58 SER C       -10.61       -25.37       67.6         58 SER C       -10.61       -25.37       67.6         58 SER C       -9.08       -24.27       70.8         58 SER C       -9.08       -24.27       70.8         58 SER C       -10.61       -25.37       67.6         58 SER C       -10.61       -25.37       67.6         58 SER C       -10.61       -25.37       67.6         59 GLU C       -11.57       -26.29       67.5         59 GLU CA       -11.71       -27.13       66.3         59 GLU CB       -12.49       -28.41       66.7         59 GLU CC       -11.81       -29.30       67.7     <	15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00
57 VAL C -12.91 -24.19 69.15 57 VAL O -13.08 -24.04 67.95 58 SER N -11.71 -24.46 69.75 58 SER CA -10.50 -24.55 68.85 58 SER CB -9.34 -25.09 69.75 58 SER CB -9.08 -24.27 70.85 58 SER C -10.61 -25.37 67.65 58 SER C -10.61 -25.37 67.65 59 GLU N -11.57 -26.29 67.55 59 GLU CA -11.71 -27.13 66.3 59 GLU CB -12.49 -28.41 66.7 59 GLU CB -11.81 -29.30 67.7 59 GLU CD -11.90 -28.75 69.1 59 GLU CD -11.90 -28.75 69.1 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CB -14.99 -24.01 65.0 60 ASN CD -15.62 -26.23 64.3 60 ASN CD -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CB -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00
57 VAL O	15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00
58 SER N -11.71 -24.46 69.7 58 SER CA -10.50 -24.55 68.8 58 SER CB -9.34 -25.09 69.7 58 SER OG -9.08 -24.27 70.8 58 SER C -10.61 -25.37 67.6 58 SER C -10.61 -25.37 67.6 58 SER O -9.84 -25.18 66.7 59 GLU N -11.57 -26.29 67.5 59 GLU CB -12.49 -28.41 66.7 59 GLU CB -12.49 -28.41 66.7 59 GLU CD -11.90 -28.75 69.1 59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CB -14.99 -24.01 65.0 60 ASN OD1 -17.03 -25.09 65.6 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00
58 SER CA	15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00
58 SER CB	15.00 15.00 15.00 0 15.00 6 15.00 8 15.00 3 15.00 5 15.00 6 15.00
58 SER CB	15.00 15.00 15.00 0 15.00 6 15.00 8 15.00 3 15.00 5 15.00 6 15.00
58 SER C	5 15.00 2 15.00 0 15.00 6 15.00 8 15.00 3 15.00 5 15.00 6 15.00
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59 GLU N -11.57 -26.29 67.5 59 GLU CA -11.71 -27.13 66.3 59 GLU CB -12.49 -28.41 66.7 59 GLU CG -11.81 -29.30 67.7 59 GLU CD -11.90 -28.75 69.1 59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN OD1 -17.03 -25.09 65.6 60 ASN C -12.75 -23.19 64.2 60 ASN C -12.75 -23.19 64.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -22.41 63.2 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	0 15.00 6 15.00 8 15.00 3 15.00 5 15.00 6 15.00
59 GLU CA -11.71 -27.13 66.3 59 GLU CB -12.49 -28.41 66.7 59 GLU CG -11.81 -29.30 67.7 59 GLU CD -11.90 -28.75 69.1 59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	6 15.00 8 15.00 3 15.00 5 15.00 6 15.00
59 GLU CB -12.49 -28.41 66.7 59 GLU CG -11.81 -29.30 67.7 59 GLU CD -11.90 -28.75 69.1 59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN OD1 -17.03 -25.09 65.6 60 ASN OD -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	3 15.00 5 15.00 6 15.00
59 GLU CG -11.81 -29.30 67.7 59 GLU CD -11.90 -28.75 69.1 59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	5 15.00 6 15.00
59 GLU CD -11.90 -28.75 69.1 59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	6 15.00
59 GLU OE1 -12.96 -28.18 69.5 59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	
59 GLU OE2 -10.92 -28.91 69.9 59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	1 15 00
59 GLU C -12.39 -26.41 65.2 59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	¥ 13.00
59 GLU O -12.46 -26.91 64.1 60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	1 15.00
60 ASN N -12.93 -25.24 65.5 60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	3 15.00
60 ASN CA -13.61 -24.41 64.5 60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	1 15.00
60 ASN CB -14.99 -24.01 65.0 60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	3 15.00
60 ASN CG -15.97 -25.16 65.0 60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	3 15.00
60 ASN OD1 -17.03 -25.09 65.6 60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	3 15.00
60 ASN ND2 -15.62 -26.23 64.3 60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	2 15.00
60 ASN C -12.75 -23.19 64.2 60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	3 15.00
60 ASN O -11.79 -22.94 64.9 61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	1 15.00
61 ASP N -13.12 -22.41 63.2 61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	7 15.00
61 ASP CA -12.36 -21.23 62.8 61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	8 15.00
61 ASP CB -12.21 -21.20 61.3 61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	5 15.00
61 ASP CG -10.99 -20.43 60.9 61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	8 15.00
61 ASP OD1 -10.38 -19.70 61.7 61 ASP OD2 -10.63 -20.55 59.7	5 15.00
61 ASP OD2 -10.63 -20.55 59.7	0 15.00
	1 15.00
	1 15.00
61 ASP C -12.92 -19.89 63.3	9 15.00
61 ASP O -12.71 -18.86 62.7	6 15.00
62 GLY N -13.61 -19.89 64.5	2 15.00
62 GLY CA -14.16 -18.66 65.0	6 15.00
62 GLY C -15.11 -17.97 64.1	15.00
62 GLY O -16.17 -18.50 63.7	9 15.00
63 CYS N -14.73 -16.79 63.6	2 15.00
63 CYS CA -15.56 -16.06 62.6	
63 CYS C -15.39 -16.55 61.2	7 15.00
63 CYS O -16.01 -16.03 60.3	

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63 CYS CI	-15.28	-14.56	62.73	15.00
63 CYS SO	-15.94	-13.72	64.20	15.00
64 GLY N	-14.52	-17.54	61.06	15.00
64 GLY CA		-18.09	59.74	15.00
64 GLY C	-15.24	-19.23	59.47	15.00
64 GLY O	-15.42	-19.63	58.32	15.00
65 GLY N	-15.85	-19.77	60.52	15.00
65 GLY CA	-16.79	-20.86	60.33	15.00
65 GLY C	-16.54	-22.03	61.25	15.00
65 GLY O	-15.56	-22.04	62.00	15.00
66 GLY N	-17.42	-23.02	61.19	15.00
66 GLY CA	-17.29	-24.19	62.03	15.00
66 GLY C	-18.50	-25.10	61.96	15.00
66 GLY O	-19.48	-24.76	61.32	15.00
67 TYR N	-18.43	-26.25	62.62	15.00
67 TYR CA	-19.53	-27.20	62.62	15.00
67 TYR CB	-19.02	-28.60	62.32	15.00
67 TYR CG	-18.35	-28.77	60.99	15.00
67 TYR CD	1 -19.08	-29.16	59.86	15.00
67 TYR CE	1 -18.46	-29.38	58.65	15.00
67 TYR CD	2 -16.98	-28.59	60.86	15.00
67 TYR CE	2 -16.36	-28.81	59.65	15.00
67 TYR CZ	-17.10	-29.20	58.55	15.00
67 TYR OH	-16.46	-29.41	57.35	15.00
67 TYR C	-20.23	-27.22	63.97	15.00
67 TYR 0	-19.59	-27.04	65.00	15.00
68 MET N	-21.52	-27.51	63.96	15.00
68 MET CA	-22.31	-27.57	65.19	15.00
68 MET CB	-23.81	-27.69	64.91	15.00
68 MET CG	-24.46	-26.48	64.23	15.00
68 MET SD	-24.10	-26.27	62.47	15.00
68 MET CE	-25.07	-27.56	61.75	15.00
68 MET C	-21.86	-28.72	66.09	15.00
68 MET O	-21.76	-28.56	67.30	15.00
69 THR N	-21.54	-29.86	65.49	15.00
69 THR CA	-21.10	-31.02	66.26	15.00
69 THR CB	-20.78	-32.22	65.35	15.00
69 THR OG	l -20.01	-31.77	64.24	15.00
69 THR CG2	2 -22.06	-32.86	64.85	15.00
69 THR C	-19.88	-30.71	67.11	15.00
69 THR O	-19.77	-31.19	68.25	15.00
70 ASN N	-18.97	-29.89	66.59	15.00
70 ASN CA	-17.77	-29.52	67.33	15.00

70	ASN	СВ	-16.79	-28.76	66.46	15.00
70	ASN	CG	-15.98	-29.65	65.58	15.00
70	ASN	OD1	-15.42	-29.19	64.60	15.00
70	ASN	ND2	-15.89	-30.93	65.92	15.00
70	ASN	С	-18.11	-28.66	68.55	15.00
70	ASN	0	-17.46	-28.77	69.59	15.00
71	ALA	N	-19.12	-27.80	68.40	15.00
71	ALA	CA	-19.57	-26.91	69.47	15.00
71	ALA	CB	-20.58	-25.91	68.94	15.00
71	ALA	С	-20.15	-27.71	70.63	15.00
71	ALA	0	-19.80	-27.50	71.78	15.00
72	PHE	N	-21.03	-28.66	70.31	15.00
72	PHE	CA	-21.64	-29.51	71.33	15.00
72	PHE	CB	-22.57	-30.54	70.69	15.00
72	PHE	CG	-23.72	-29.93	69.96	15.00
72	PHE	CD1	-24.28	-28.73	70.39	15.00
72	PHE	CD2	-24.24	-30.55	68.84	15.00
72	PHE	CE1	-25.33	-28.16	69.70	15.00
72	PHE	CE2	-25.30	-29.98	68.15	15.00
72	PHE	CZ	-25.84	-28.78	68.58	15.00
72	PHE	C	-20.53	-30.25	72.07	15.00
72	PHE	0	-20.43	-30.18	73.31	15.00
73	GLN	N	-19.67	-30.90	71.31	15.00
73	GLN	CA	-18.56	-31.66	71.86	15.00
73	GLN	CB	-17.68	-32.20	70.72	15.00
73	GLN	CG	-16.78	-33.40	71.09	15.00
73	GLN	CD	-17.54	-34.73	71.21	15.00
73	GLN	OE1	-17.47	-35.58	70.31	15.00
73	GLN	NE2	-18.21	-34.94	72.35	15.00
73	GLN	С	-17.76	-30.78	72.84	15.00
73	GLN	0	-17.33	-31.25	73.89	15.00
74	TYR	N	-17.62	-29.50	72.53	15.00
74	TYR	CA	-16.89	-28.59	73.40	15.00
74	TYR	CB	-16.70	-27.22	72.75	15.00
74	TYR	CG	-16.35	-26.13	73.74	15.00
74	TYR	CD1	-15.09	-26.05	74.30	15.00
74	TYR	CE1	-14.77	-25.07	75.22	15.00
74	TYR	CD2	-17.30	-25.18	74.12	15.00
74	TYR	CE2	-17.00	-24.19	75.04	15.00
74	TYR	CZ	-15.73	-24.14	75.59	15.00
74	TYR	OH	-15.42	-23.15	76.51	15.00
74	TYR	С	-17.58	-28.38	74.73	15.00
74	TYR	0	-16.93	-28.39	75.78	15.00

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	VAI		-18.88	-28.12	74.67	15.00
75		CA	-19.68	-27.89	75.88	15.00
75		CB	-21.15	-27.58	75.52	15.00
75	VAI	CG1	-21.95	-27.27	76.78	15.00
75	VAL	CG2	-21.22	-26.39	74.55	15.00
75	VAL	C	-19.62	-29.10	76.80	15.00
75	VAL	. 0	-19.60	-28.96	78.03	15.00
76	GLN	N	-19.59	-30.29	76.21	15.00
76	GLN	CA	-19.51	-31.51	76.98	15.00
76	GLN	CB	-19.75	-32.69	76.05	15.00
76	GLN	CG	-19.79	-34.05	76.69	15.00
76	GLN	CD	-19.56	-35.13	75.66	15.00
76	GLN	OE1	-20.03	-35.04	74.53	15.00
76	GLN	NE2	-18.78	-36.13	76.03	15.00
76	GLN	С	-18.14	-31.60	77.64	15.00
76	GLN	0	-18.03	-31.59	78.86	15.00
77	LYS	N	-17.08	-31.61	76.84	15.00
77	LYS	CA	-15.72	-31.70	77.36	15.00
77	LYS	CB	-14.70	-31.55	76.22	15.00
77	LYS	CG	-13.27	-31.34	76.69	15.00
77	LYS	CD	-12.32	-31.10	75.51	15.00
77	LYS	CE	-10.89	-30.81	75.97	15.00
77	LYS	NZ	-10.30	-31.95	76.76	15.00
77	LYS	С	-15.45	-30.64	78.42	15.00
77	LYS	0	-14.81	-30.91	79.45	15.00
78	ASN	N	-15.92	-29.42	78.17	15.00
78	ASN	CA	-15.74	-28.29	79.06	15.00
78	ASN		-15.98	-27.00	78.28	15.00
78	ASN	CG	-15.69	-25.76	79.10	15.00
78	ASN	OD1	-14.58	-25.59	79.62	15.00
78	ASN	ND2	-16.67	-24.87	79.19	15.00
78	ASN	C	-16.68	-28.34	80.25	15.00
78	ASN	0	-16.42	-27.72	81.28	15.00
79	ARG		-17.79	-29.06	80.11	15.00
79	ARG		-18.78	-29.18	81.16	15.00
79	ARG	CB	-18.14	-29.76	82.43	15.00
79	ARG	CG	-17.67	-31.20	82.26	15.00
79	ARG	CD	-16.65	-31.59	83.33	15.00
79	ARG	NE	-17.15	-31.38	84.68	15.00
79	ARG	CZ	-18.16	-32.06	85.22	15.00
79	ARG	NH1	-18.78	-33.03	84.55	15.00
79	ARG		-18.62	-31.70	86.43	15.00
79	ARG	С	-19.45	-27.84	81.45	15.00

79	ARG	0	-19.31	-27.30	82.55	15.00
80	GLY	N	-20.13	-27.28	80.45	15.00
80	GLY	CA	-20.82	-26.02	80.66	15.00
80	GLY	С	-20.62	-24.91	79.64	15.00
80	GLY	0	-19.56	-24.82	79.00	15.00
81	ILE	N	-21.64	-24.08	79.48	15.00
81	ILE	CA	-21.59	-22.95	78.57	15.00
81	ILE	СВ	-22.30	-23.23	77.22	15.00
81	ILE	CG2	-23.77	-23.52	77.45	15.00
81	ILE	CG1	-22.10	-22.05	76.27	15.00
81	ILE	CD1	-22.84	-22.16	74.96	15.00
81	ILE	С	-22.24	-21.75	79.25	15.00
81	ILE	0	-23.28	-21.89	79.90	15.00
82	ASP	N	-21.61	-20.59	79.11	15.00
82	ASP	CA	-22.11	-19.37	79.71	15.00
82	ASP	CB	-21.03	-18.29	79.64	15.00
82	ASP	CG	-19.90	-18.53	80.58	15.00
82	ASP	OD1	-18.82	-17.98	80.34	15.00
82	ASP	OD2	-20.09	-19.26	81.58	15.00
82	ASP	C	-23.36	-18.81	79.09	15.00
82	ASP	0	-23.69	-19.07	77.93	15.00
83	SER	N	-24.07	-18.02	79.89	15.00
83	SER	CA	-25.27	-17.36	79.44	15.00
83	SER	CB	-26.09	-16.90	80.64	15.00
83	SER	OG	-25.27	-16.23	81.59	15.00
83	SER	С	-24.75	-16.15	78.66	15.00
83	SER	0	-23.57	-15.79	78.79	15.00
84	GLU	N	-25.61	-15.54	77.86	15.00
84	GLU	CA	-25.25	-14.36	77.07	15.00
84	GLU	CB	-26.46	-13.84	76.31	15.00
84	GLU	CG	-26.17	-12.64	75.43	15.00
84	GLU	CD	-25.31	-12.99	74.24	15.00
84	GLU	OEl	-24.08	-12.98	74.38	15.00
84	GLU	OE2	-25.87	-13.29	73.17	15.00
84	GLU	С	-24.70	-13.27	77.99	15.00
84	GLU	0	-23.53	-12.89	77.88	15.00
85	ASP	N	-25.51	-12.82	78.94	15.00
85	ASP	CA	-25.09	-11.79	79.87	15.00
85	ASP	CB	-26.13	-11.60	80.99	15.00
85	ASP	CG	-25.66	-10.63	82.08	15.00
85	ASP	OD1	-25.92	-10.88	83.29	15.00
85	ASP	OD2	-25.03	-9.60	81.73	15.00
85	ASP	С	-23.72	-12.09	80.49	15.00

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	SP O	-22.91	-11.18	80.65	15.00
	LA N	-23.45	-13.34	80.81	15.00
	LA CA	-22.18	-13.67	81.44	15.00
86 A	LA CB	-22.25	-15.05	82.03	15.00
86 A	LA C	-21.01	-13.56	80.47	15.00
86 A	LA O	-19.91	-13.17	80.86	15.00
87 T	YR N	-21.26	-13.89	79.21	15.00
87 T	YR CA	-20.23	-13.85	78.18	15.00
87 T	YR CB	-19.77	-15.27	77.87	15.00
87 T	YR CG	-18.42	-15.39	77.19	15.00
87 T	YR CD1	-17.96	-14.41	76.30	15.00
87 T	YR CE1	-16.74	-14.56	75.65	15.00
87 T	YR CD2	-17.62	-16.52	77.40	15.00
87 T	CR CE2	-16.40	-16.67	76.76	15.00
87 T	R CZ	-15.96	-15.69	75.88	15.00
87 m	CR OH	-14.75	-15.83	75.25	15.00
	CR C	-20.93	-13.23	76.97	15.00
	TR O	-21.57	-13.94	76.19	15.00
	RO N	-20.90	-11.90	76.86	15.00
	O CD	-20.42	-10.97	77.90	15.00
	O CA	-21.52	-11.15	75.78	15.00
	O CB	-21.33	-9.70	76.23	15.00
	O CG	-21.34	-9.81	77.71	15.00
	o c	-20.91	-11.38	74.39	15.00
	0 0	-19.74	-11.74	74.25	15.00
	RN	-21.73	-11.12	73.38	15.00
	R CA	-21.35	-11.29	71.99	15.00
	R CB	-22.56	-11.78	71.18	15.00
	R CG	-22.24	-12.15	69.76	15.00
	R CD1	-21.38	-13.20	69.47	15.00
	R CE1	-21.06	-13.53	68.17	15.00
	R CD2	-22.78	-11.44	68.70	15.00
	R CE2	-22.47	-11.76	67.39	15.00
89 TY		-21.61	-12.79	67.13	15.00
	R OH	-21.27	-13.08	65.83	15.00
	RC	-20.85	-9.95	71.48	15.00
	RO	-21.52	-8.92	71.64	15.00
	LN	-19.65	-9.94	70.91	15.00
	L CA	-19.07	-8.71	70.39	15.00
	L CB	-17.75	-8.36	71.12	15.00
	L CG1	-17.97	-8.36	72.62	15.00
	L CG2	-16.65	-ÿ.33	70.74	15.00
90 VA	r c	-18.88	-8.78	68.87	15.00

90	VAL	0	-18.54	-7.78	68.23	15.00
91	GLY	N	-19.08	-9.96	68.30	15.00
91	GLY	CA	-18.95	-10.12	66.86	15.00
91	GLY	С	-17.55	-10.23	66.27	15.00
91	GLY	0	-17.34	-9.87	65.12	15.00
92	GLN	N	-16.61	-10.80	67.01	15.00
92	GLN	CA	-15.24	-10.95	66.50	15.00
92	GLN	CB	-14.56	-9.58	66.41	15.00
92	GLN	CG	-14.68	-8.72	67.67	15.00
92	GLN	CD	-13.59	-7.67	67.79	15.00
92	GLN	OE1	-12.77	-7.72	68.72	15.00
92	GLN	NE2	-13.56	-6.72	66.85	15.00
92	GLN	С	-14.45	-11.86	67.43	15.00
92	GLN	0	-14.78	-11.97	68.62	15.00
93	GLU	N	-13.43	-12.52	66.89	15.00
93	GLU	CA	-12.64	-13.42	67.71	15.00
93	GLU	CB	-11.68	-14.28	66.89	15.00
93	GLU	CG	-12.31	-15.10	65.78	15.00
93	GLU	CD	-11.63	-14.87	64.44	15.00
93	GLU	OE1	-11.94	-15.61	63.48	15.00
93	GLU	OE2	-10.78	-13.94	64.33	15.00
93	GLU	С	-11.84	-12.62	68.71	15.00
93	GLU	0	-11.41	-11.49	68.44	15.00
94	GLU	N	-11.61	-13.26	69.85	15.00
94	GLU	CA	-10.88	-12.70	70.98	15.00
94	GLU	CB	-11.81	-11.81	71.79	15.00
94	GLU	CG	-13.19	-12.42	71.93	15.00
94	GLU	CD	-14.06	-11.70	72.92	15.00
94	GLU	OE1	-13.99	-10.45	72.96	15.00
94	GLU	OE2	-14.83	-12.38	73.64	15.00
94	GLU	С	-10.52	-13.92	71.80	15.00
94	GLU	0	-10.89	-15.02	71.45	15.00
95	SER	N	-9.81	-13.73	72.91	15.00
95	SER	CA	-9.43	-14.85	73.75	15.00
95	SER	CB	-8.32	-14.42	74.71	15.00
95	SER	OG	-7.20	-13.94	73.98	15.00
95	SER	C	-10.62	-15.42	74.52	15.00
95	SER	0	-11.48	-14.67	75.02	15.00
96	CYS	N	-10.69	-16.75	74.57	15.00
96	CYS	CA	-11.76	-17.45	75.28	15.00
96	CYS	C	-11.74	-16.96	76.71	15.00
96	CYS	0	-10.73	-17.09	77.42	15.00
96	CYS	CB	-11.53	-18.97	75.20	15.00

			**	
96 CYS SG	-12.62	-20.03	76.22	15.00
97 MET N	-12.85	-16.38	77.14	15.00
97 MET CA	-12.96	-15.85	78.49	15.00
97 MET CB	-13.32		78.39	15.00
97 MET CG	-12.29		77.69	15.00
97 MET SD	-13.01		77.03	15.00
97 MET CE	-14.07	-11.44	78.40	15.00
97 MET C	-14.03	-16.60	79.28	15.00
97 MET O	-14.87	-15.99	79.96	15.00
98 TYR N	-13.98	-17.93	79.21	15.00
98 TYR CA	-14.96	-18.74	79.91	15.00
98 TYR CB	-14.69	-20.23	79.71	15.00
98 TYR CG	-15.74	-21.10	80.34	15.00
98 TYR CD1	-17.08		79.97	15.00
98 TYR CE1		-21.74	80.59	15.00
98 TYR CD2	-15.41	-22.00	81.36	15.00
98 TYR CE2	-16.40	-22.76	81.98	15.00
98 TYR CZ	-17.72	-22.62	81.60	15.00
98 TYR OH	-18.70	-23.34	82.23	15.00
98 TYR C	-15.03	-18.43	81.39	15.00
98 TYR O	-14.01	-18.38	82.08	15.00
99 ASN N	-16.25	-18.27	81.88	15.00
99 ASN CA	-16.49	-17.97	83.28	15.00
99 ASN CB	-17.29	-16.66	83.38	15.00
99 ASN CG	-17.66	-16.33	84.80	15.00
99 ASN OD1	-16.88	-16.57	85.74	15.00
99 ASN ND2	-18.85	-15.79	84.99	15.00
99 ASN C	-17.28	-19.10	83.92	15.00
99 ASN O	-18.51	-19.16	83.78	15.00
100 PRO N	-16.60	-19.99	84.67	15.00
100 PRO CD	-15.22	-19.88	85.16	15.00
100 PRO CA	-17.29	-21.11	85.32	15.00
100 PRO CB	-16.20	-21.72	86.20	15.00
100 PRO CG	-15.31	-20.55	86.51	15.00
100 PRO C		-20.63	86.16	15.00
100 PRO O	-19.51	-21.24	86.15	15.00
101 THR N		-19.50	86.84	15.00
101 THR CA	-19.33	-18.95	87.68	15.00
101 THR CB		-17.68	88.45	15.00
101 THR OG1		-16.57	87.55	15.00
101 THR CG2		-17.93	89.13	15.00
101 THR C		-18.59	86.82	15.00
101 THR O	-21.68	-18.56	37.32	15.00

102	GLY	N	-20.32	-18.34	85.54	15.00
102	GLY	CA	-21.40	-17.98	84.64	15.00
102	GLY	С	-22.06	-19.12	83.88	15.00
102	GLY	0	-22.92	-18.86	83.03	15.00
103	LYS	N	-21.65	-20.36	84.12	15.00
103	LYS	CA	-22.24	-21.50	83.42	15.00
103	LYS	СВ	-21.72	-22.83	83.98	15.00
103	LYS	CG	-22.32	-24.05	83.29	15.00
103	LYS	CD	-22.10	-25.33	84.06	15.00
103	LYS	CE	-22.96	-25.41	85.35	15.00
103	LYS	NZ	-24.41	-25.79	85.14	15.00
103	LYS	С	-23.75	-21.49	83.57	15.00
103	LYS	0	-24.26	-21.26	84.67	15.00
104	ALA	N	-24.47	-21.73	82.48	15.00
104	ALA	CA	-25.93	-21.75	82.53	15.00
104	ALA	CB	-26.51	-20.51	81.87	15.00
104	ALA	С	-26.52	-22.99	81.89	15.00
104	ALA	0	-27.73	-23.15	81.87	15.00
105	ALA	N	-25.66	-23.87	81.38	15.00
105	ALA	CA	-26.11	-25.10	80.75	15.00
105	ALA	CB	-26.84	-24.78	79.44	15.00
105	ALA	С	-24.95	-26.03	80.46	15.00
105	ALA	0	-23.79	-25.67	80.62	15.00
106	LYS	N	-25.28	-27.26	80.07	15.00
106	LYS	CA	-24.29	-28.25	79.70	15.00
106	LYS	CB	-23.55	-28.80	80.92	15.00
106	LYS	CG	-24.41	-29.35	82.04	15.00
106	LYS	CD	-23.54	-29.64	83.27	15.00
106	LYS	CE	-22.40	-30.62	82.95	15.00
106	LYS	NZ	-21.34	-30.67	84.03	15.00
106	LYS	С	-24.99	-29.36	78.93	15.00
106	LYS		-26.21	-29.30	78.75	15.00
	CYS		-24.23	-30.30	78.39	15.00
107	CYS	CA		-31.40	77.64	15.00
	CYS		-25.06		76.19	15.00
107	CYS	SG		-30.95	75.18	15.00
107	CYS	С	-23.91	-32 61	77.68	15.00
107	CYS	0	-22.75	-32.49	78.06	15.00
	ARG		-24.43	-33.77	77.32	15.00
	ARG			-35.00	77.33	1.5.00
	ARG		-24.12		78.42	15.00
108	ARG	CG		-35.03	78.64	15.00
108	ARG	CD	-26.27	-37.28	78.04	15.00

108	ARG	NE	-27.73	-37.21	78.14	15.00
	ARG	CZ	-28.57	-38.06	77.54	15.00
108	ARG	NH1	-28.09	-39.05	76.80	15.00
108	ARG	NH2	-29.88	-37.94	77.72	15.00
108	ARG	C	-23.55	-35.70	75.97	15.00
108	ARG	0	-23.77	-36.91	75.85	15.00
109	GLY	N	-23.19	-34.93	74.94	15.00
109	GLY	CA	-23.06	-35.50	73.61	15.00
109	GLY	С	-24.08	-34.95	72.65	15.00
109	GLY	0	-24.81	-34.01	72.97	15.00
110	TYR	N	-24.16	-35.57	71.48	15.00
110	TYR	CA	-25.07	-35.16	70.42	15.00
110	TYR	CB	-24.41	-34.09	69.55	15.00
110	TYR	CG	-23.10	-34.53	68.92	15.00
110	TYR	CD1	-21.91	-34.49	69.64	15.00
110	TYR	CE1	-20.71	-34.90	69.08	15.00
110	TYR	CD2	-23.06	-34.99	67.62	15.00
110	TYR	CE2	-21.87	-35.41	67.04	15.00
	TYR	CZ	-20.70	-35.35	67.77	15.00
110	TYR	OH	-19.52	-35.75	67.18	15.00
110	TYR	С	-25.39	-36.37	69.57	15.00
110	TYR	0	-24.80	-37.44	69.77	15.00
111	ARG	N	-26.29	-36.20	68.61	15.00
111	ARG		-26.69	-37.28	67.73	15.00
	ARG	CB	-27.96	-37.95	68.26	15.00
111	ARG	CG	-27.84	-38.48	69.67	15.00
	ARG	CD	-29.18	-39.01	70.20	15.00
111	ARG	NE	-29.77	-40.08	69.39	15.00
	ARG	CZ	-29.16	-41.20	69.02	15.00
	ARG	NH1	-29.81	-42.09	68.27	15.00
111	ARG		-27.91	-41.45	69.40	15.00
	ARG		-26.98	-36.73	66.35	15.00
	ARG		-27.99	-36.05	66.17	15.00
	GLU			-36.96	65.39	15.00
•	GLU		-26.32	-36.48	64.03	15.00
	GLU		-25.09	-36.71	63.15	15.00
	GLU		-23.91	-35.81	63.46	15.00
	GLU		-23.40	-35.09	62.22	15.00
	GLU		-24.13	-34.21	61.69	15.00
	GLU		-22.27	-35.41	61.77	15.00
		С		-37.23	63.45	15.00
	GLU			-38.41	63.76	15.00
113	ILE	N	-28.29	-36.55	62.64	15.00

113	ILE	CA	-29.46	-37.15	62.02	15.00
113	ILE	CB	-30.51	-36.04	61.71	15.00
113	ILE	CG2	-31.75	-36.60	61.05	15.00
113	ILE	CG1	-30.92	-35.38	63.02	15.00
113	ILE	CD1	-31.95	-34.32	62.87	15.00
113	ILE	С	-28.95	-37.84	60.75	15.00
113	ILE	0	-27.93	-37.44	60.20	15.00
114	PRO	N	-29.60	-38.95	60.34	15.00
114	PRO	CD	-30.69	-39.67	61.02	15.00
114	PRO	CA	-29.17	-39.68	59.14	15.00
114	PRO	CB	-30.28	-40.72	58.97	15.00
114	PRO	CG	-30.62	-41.05	60.38	15.00
114	PRO	С	-29.04	-38.79	57.93	15.00
114	PRO	0	-30.00	-38.17	57.47	15.00
115	GLU	N	-27.82	-38.75	57.41	15.00
115	GLU	CA	-27.50	-37.92	56.26	15.00
115	GLU		-26.12	-38.30	55.74	15.00
	GLU		-25.58	-37.36	54.68	15.00
	GLU	CD	-24.19	-37.76	54.22	15.00
115			-23.20	-37.34	54.86	15.00
115	GLU		-24.10	-38.51	53.22	15.00
115	GLU		-28.52	-38.00	55.14	15.00
	GLU		-28.72	-39.05	54.56	15.00
	GLY		-29.21	-36.89	54.90	15.00
	GLY		-30.18	-36.81	53.83	15.00
	GLY		-31.55	-37.41	54.07	15.00
116			-32.34	-37.53	53.14	15.00
	ASN		-31.86	-37.73	55.32	15.00
	ASN		-33.15	-38.34	55.65	15.00
117			-32.91	-39.54	56.56	15.00
117			-34.17	-40.32	56.84	15.00
117			-35.26	-39.75	56.98	15.00
	ASN		-34.04	-41.64	56.94	15.00
	ASN		-34.11			15.00
	ASN		-34.16		57.52	15.00
	GLU			-36.66	55.48	15.00
	GLU			-35.67	55.98	15.00
	GLU		-36.67	-35.08	54.85	15.00
	GLU		-35.91	~34.08	54.01	15.00
	GLU			-32.98	53.50	15.00
	GLU			-32.38	54.32	15.00
				-32.73	52.29	15.00
118	GLU	C	-36.80	-36.20	57.04	15.00

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	B GLU	0	-37.05	-35.54	58.04	15.00
119		N	-37.34	-37.38	56.81	15.00
119		CA	-38.28	-37.95	57. <b>7</b> 7	15.00
119		CB	-38.87	-39.24	57.21	15.00
119		CG	-39.46	-39.06	55.80	15.00
119		CD	-40.57	-38.01	55.77	15.00
119	LYS	CE	-41.82	-38.49	56.49	15.00
119	LYS	NZ	-42.97	-37.55	56.32	15.00
119	LYS		-37.66	-38.15	59.15	15.00
119	LYS	0	-38.29	-37.87	60.16	15.00
120	ALA	N	-36.39	-38.56	59.19	15.00
120	ALA	CA	-35.73	-38.76	60.48	15.00
120	ALA	CB	-34.39	-39.43	60.30	15.00
120			-35.56	-37.41	61.15	15.00
120	ALA	0	-35.40	-37.34	62.37	15.00
121	LEU	N	-35.58	-36.35	60.34	15.00
121			-35.45	-34.99	60.83	15.00
121	LEU		-35.03	-34.03	59.71	15.00
121	LEU	CG	-34.92	-32.51	59.96	15.00
121		CD1	-33.98	-32.21	61.11	15.00
121			-34.45	-31.82	58.71	15.00
121		С	-36.78	-34.54	61.43	15.00
121	LEU (	0	-36.80	-33.96	62.51	15.00
122	LYS		-37.89	-34.84	60.76	15.00
122	LYS	CA	-39.20	-34.44	61.28	15.00
122	LYS (	CB	-40.34	-34.86	60.35	15.00
122	LYS	CG	-41.71	-34.49	60.95	15.00
122	LYS (	CD	-42.90	-34.72	60.02	15.00
122	LYS (	CE	-43.21	-36.19	59.84	15.00
122	LYS 1	NZ	-42.13	-36.87	59.07	15.00
122	LYS (	С	-39.43	-35.03	62.67	15.00
122	LYS (		-40.00	-34.38	63.54	15.00
123	ARG 1		-39.02	-36.28	62.85	15.00
123	ARG (	CA	-39.18	-36.96	64.12	15.00
123			-38.90	-38.45	63.95	15.00
123			-40.04	-39.22	63.30	15.00
123		CD	-39.53	-40.52	62.67	15.00
123			-38.42	-41.08	63.44	15.00
	ARG C			-41.85	62.92	15.00
123				-42.16	61.62	15.00
	ARG 1		-36.45	-42.23	63.68	15.00
	ARG C			-36.34	65.14	15.00
123	ARG C	) '	-38.65	-36.04	66.25	15.00

124	ALA	N	-36.99	-36.12	64.76	15.00
124	ALA	CA	-36.05	-35.51	65.68	15.00
124	ALA	CB	-34.70	-35.31	65.02	15.00
124	ALA	С	-36.60	-34.17	66.19	15.00
124	ALA	0	-36.55	-33.91	67.39	15.00
125	VAL	N	-37.14	-33.34	65.30	15.00
125	VAL	CA	-37.68	-32.06	65.76	15.00
125	VAL	CB	-38.01	-31.03	64.60	15.00
125	VAL	CG1	-36.78	-30.27	64.19	15.00
125	VAL	CG2	-38.58	-31.72	63.39	15.00
125	VAL	С	-38.94	-32.28	66.58	15.00
125	VAL	0	-39.21	-31.52	67.50	15.00
126	ALA	N	-39.69	-33.32	66.27	15.00
126	ALA	CA	-40.93	-33.60	66.98	15.00
126	ALA	СВ	-41.81	-34.52	66.16	15.00
126	ALA	С	-40.75	-34.16	68.38	15.00
126	ALA	0	-41.53	-33.84	69.28	15.00
127	ARG	N	-39.73	-35.00	68.55	15.00
127	ARG	CA	-39.42	-35.68	69.81	15.00
127	ARG	CB	-39.04	-37.14	69.54	15.00
127	ARG	CG	-40.20	-37.95	69.00	15.00
127	ARG	CD	-39.78	-39.23	68.27	15.00
127	ARG	NE	-40.95	-39.80	67.60	15.00
127	ARG	CZ	-40.95	-40.88	66.83	15.00
127	ARG	NH1	-42.09	-41.30	66.28	15.00
127	ARG	NH2	-39.84	-41.56	66.63	15.00
127	ARG	С	-38.33	-35.04	70.66	15.00
127	ARG	0	-38.25	-35.30	71.86	15.00
128	VAL	N	-37.48	-34.22	70.04	15.00
128	VAL	CA	-36.40	-33.54	70.75	15.00
128	VAL	CB	-35.03	-33.81	70.10	15.00
128	VAL	CG1	-33.92	-33.34	71.02	15.00
128	VAL	CG2	-34.87	-35.29	69.78	15.00
128	VAL	С	-36.58	-32.02	70.88	15.00
128	VAL	0	-36.43	-31.46	71.95	15.00
129	GLY	N	-36.89	-31.35	69.77	1.5.00
129	GLY	CA	-37.08	-29.91	69.81	15.00
129	GLY	C	-36.26	-29.29	68.69	15.00
129	GLY	0	-36.02	-29.36	67.68	15.00
130	PRO	N	-35.83	-28.02	68.81	15.00
130	PRO	CD	-36.20	-27.06	69.86	15.00
1.30	PRO	CA	-35.04	-27.37	67.77	15.00
130	PRO	СВ	-34.67	-26.05	68.43	15.00

130	PRO	CG	-35.92	-25.74	69.18	15.00
130	PRO	С	-33.81	-28.17	67.39	15.00
130	PRO	0	-33.07	-28.64	68.26	15.00
131	VAL	N	-33.60	-28.33	66.09	15.00
131	VAL	CA	-32.46	-29.08	65.58	15.00
131	VAL	CB	-32.94	-30.26	64.68	15.00
131	VAL	CG1	-31.76	-31.03	64.14	15.00
131	VAL	CG2	-33.82	-31.20	65.46	15.00
131	VAL	С	-31.50	-28.20	64.77	15.00
131	VAL	0	-31.93	-27.35	63.99	15.00
132	SER	N	-30.20	-28.39	64.96	15.00
132	SER	CA	-29.23	-27.61	64.22	15.00
132	SER	CB	-27.88	-27.61	64.94	15.00
132	SER	OG	-28.00	-26.96	66.20	15.00
132	SER	С	-29.08	-28.21	62.82	15.00
132	SER	0	-28.83	-29.41	62.68	15.00
133	VAL	N	-29.31	-27.39	61.80	15.00
133	VAL	CA	-29.22	-27.82	60.40	15.00
133	VAL	CB	-30.60	-27.81	59.68	15.00
133	VAL	CG1	-31.51	-28.90	60.23	15.00
133	VAL	CG2	-31.27	-26.45	59.80	15.00
133	VAL	C	-28.26	-26.93	59.62	15.00
	VAL		-27.88	-25.85	60.08	15.00
134			-27.93	-27.36	58.41	15.00
134			-27.02	-26.64	57.54	15.00
134			-25.69	-27.35	57.48	15.00
134 .			-27.64	-26.61	56.16	15.00
134 .	ALA	0	-27.92	-27.66	55.60	15.00
	ILE		-27.84	-25.43	55.60	15.00
	ILE		-28.45	-25.31	54.28	15.00
	ILE		-29.84	-24.62	54.36	15.00
	ILE		-30.82	-25.47	55.15	15.00
	ILE		-29.70	-23.24	55.00	15.00
135						15.00
135			-27.59		53.32	15.00
135			-26.49		53.66	15.00
136			-28.09	-24.33	52.10	15.00
136				-23.52	51.07	15.00
136				-24.23	49.72	15.00
136						15.00
136				-23.40		
				-22.50		15.00
136 /	ASP	С	-28.22	-22.21	50.99	15.00

136	ASP	0	-29.36	-22.17	50.52	15.00
137	ALA	N	-27.61	-21.13	51.46	15.00
137	ALA	CA	-28.26	-19.83	51.42	15.00
137	ALA	CB	-28.42	-19.30	52.83	15.00
137	ALA	С	-27.46	-18.84	50.56	15.00
137	ALA	0	-27.34	-17.66	50.89	15.00
138	SER	N	-26.92	-19.34	49.45	15.00
138	SER	CA	-26.12	-18.53	48.53	15.00
138	SER	CB	-25.09	-19.42	47.83	15.00
138	SER	OG	-25.71	-20.52	47.19	15.00
138	SER	С	-26.97	-17.80	47.49	15.00
138	SER	0	-26.60	-16.72	47.01	15.00
139	LEU	N	-28.12	-18.36	47.17	15.00
139	LEU	CA	-29.02	-17.79	46.19	15.00
139	LEU	CB	-30.07	-18.84	45.80	15.00
139	LEU	CG	-29.49	-20.25	45.62	15.00
139	LEU	CD1	-30.58	-21.21	45.21	15.00
139	LEU	CD2	-28.37	-20.28	44.61	15.00
139	LEU	С	-29.70	-16.52	46.70	15.00
139	LEU	0	-30.06	-16.43	47.88	15.00
140	THR	N	-29.90	-15.54	45.81	15.00
140	THR	CA	-30.54	-14.29	46.18	15.00
140	THR	CB	-30.46	-13.21	45.07	15.00
140	THR		-30.85	-13.77	43.82	15.00
140	THR		-29.05	-12.66	44.96	15.00
140	THR	С	-32.00	-14.49	46.57	15.00
140	THR	0	-32.50	-13.79	47.45	15.00
141	SER	N	-32.68	-15.45	45.95	15.00
141	SER	CA	-34.08	-15.70	46.30	15.00
141	SER		-34.66	-16.86	45.50	15.00
141	SER		-33.72	-17.92	45.39	15.00
141	SER		-34.19	-15.94	47.79	15.00
141	SER		-35.04	-15.37	48.46	15.00
	PHE				48.33	15.00
	PHE			-17.01	49.76	15.00
142	PHE		-32.21		50.11	15.00
142	PHE		-32.17	-18.41	51.57	15.00
142	PHE		-32.97	-19.43	52.08	15.00
	PHE			-17.72	52.45	15.00
142	PHE			-19.75	53.43	15.00
	PHE			-18.04	53.80	15.00
	PHE			-19.05	54.29	15.00
142	PHE	С	-33.01	-15.75	50.54	15.00

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142 PHE 0	-33.69		51.52	15.00
143 GLN N	-32.01	-15.00	50.09	15.00
143 GLN CA	-31.61	_	50.78	15.00
143 GLN CB	-30.30		50.21	15.00
143 GLN CG	-29.18	-14.29	50.23	15.00
143 GLN CD	-27.85		49.81	15.00
143 GLN OE1	-27.29	-12.84	50.48	15.00
143 GLN NE2	-27.31	-14.22	48.70	15.00
143 GLN C	-32.67		50.84	15.00
143 GLN O	-32.79	-12.04	51.86	15.00
144 PHE N	-33.45	-12.51	49.78	15.00
144 PHE CA	-34.50	-11.48	49.83	15.00
144 PHE CB	-34.57	-10.60	48.55	15.00
144 PHE CG	-34.78		47.27	15.00
144 PHE CD1	-33.92	-11.14	46.19	15.00
144 PHE CD2	-35.84	. – –	47.12	15.00
144 PHE CE1	-34.11	-11.81	44.99	15.00
144 PHE CE2	-36.04	-12.92	45.91	15.00
144 PHE CZ	-35.18	_	44.85	15.00
144 PHE C	-35.88	-12.04	50.20	15.00
144 PHE O	-36.90	-11.36	50.02	15.00
145 TYR N	-35.89	-13.28	50.71	15.00
145 TYR CA	-37.12	-13.95	51.12	15.00
145 TYR CB	-36.80	-15.21	51.94	15.00
145 TYR CG	-37.98	-15.77	52.70	15.00
145 TYR CD1	-38.84	-16.69	52.12	15.00
145 TYR CE1	-39.96	-17.15	52.80	15.00
145 TYR CD2	-38.27	-15.32	53.99	15.00
145 TYR CE2	-39.40	-15.78	54.67	15.00
145 TYR CZ	-40.24	-16.69	54.07	15.00
145 TYR OH	-41.38	-17.11	54.73	15.00
145 TYR C	-37.90	-12.98	51.97	15.00
145 TYR O	-37.32		52.74	15.00
146 SER N	-39.21	-13.02	51.84	15.00
146 SER CA	-40.07	-12.13	52.59	15.00
146 SER CB	-40.63	-11.07	51.63	15.00
146 SER OG	-41.38	-10.08	52.30	15.00
146 SER C	-41.21	-12.89	53.24	15.00
146 SER O	-41.48	-12.72	54.43	15.00
147 LYS N	-41.86	-13.77	52.48	15.00
147 LYS CA	-42.98	-14.54	53.01	15.00
147 LYS CB	-44.25	-13.71	53.04	1.5.00
147 LYS CG	-44.62	-13.11	51.70	15.00

147	LYS	CD	-46.07	-12.67	51.68	15.00
147	LYS	CE	-46.47	-12.13	50.31	15.00
147	LYS	NZ	-47.97	-12.11	50.14	15.00
147	LYS	С	-43.21	-15.79	52.19	15.00
147	LYS	0	-42.55	-16.01	51.17	15.00
148	GLY	N	-44.16	-16.61	52.64	15.00
148	GLY	CA	-44.49	-17.85	51.95	15.00
148	GLY	С	-43.47	-18.95	52.16	15.00
148	GLY	0	-42.52	-18.79	52.93	15.00
149	VAL	N	-43.64	-20.04	51.43	15.00
149	VAL	CA	-42.75	-21.19	51.52	15.00
149	VAL	CB	-43.57	-22.51	51.47	15.00
149	VAL	CG1	-42.66	-23.71	51.33	15.00
149	VAL	CG2	-44.41	-22.65	52.72	15.00
149	VAL	С	-41.67	-21.17	50.43	15.00
149	VAL	0	-41.96	-21.34	49.24	15.00
150	TYR	N	-40.43	-20.96	50.84	15.00
150	TYR	CA	-39.30	-20.91	49.91	15.00
150	TYR	CB	-38.04	-20.41	50.64	15.00
150	TYR	CG	-36.82	-20.29	49.75	15.00
150	TYR	CD1	-36.78	-19.38	48.69	15.00
150	TYR	CE1	-35.67	-19.27	47.88	15.00
150	TYR	CD2	-35.69	-21.07	49.97	15.00
150	TYR	CE2	-34.56	-20.96	49.16	15.00
150	TYR	CZ	-34.56	-20.06	48.11	15.00
150	TYR	OH	-33.45	-19.93	47.32	15.00
150	TYR	С	-39.03	-22.26	49.27	15.00
150	TYR	0	-39.23	-23.31	49.88	15.00
151	TYR	N	-38.55	-22.22	48.03	15.00
151	TYR	CA	-38.21	-23.42	47.28	15.00
151	TYR	CB	-39.45	-24.26	46.99	15.00
151	TYR	CG	-39.15	-25.46	46.11	15.00
	TYR		-38.22	-26.42	46.51	15.00
151	TYR	CE1	-37.94	-27.53	45.71	15.00
151	TYR	CD2	-39.79		44.89	15.00
151	TYR	CE2	-39.52	-26.74	44.08	15.00
151	TYR	CZ	-38.59	-27.69	44.50	15.00
151	TYR	OH	-38.36	-28.83	43.75	15.00
151	TYR	С	-37.60	-23.00	45.97	15.00
151	TYR	0	-38.29	-22.44	45.11	15.00
152	ASP	N		-23.25	45.80	15.00
152	ASP	CA	-35.66	-22.90	44.55	15.00
152	ASP	CB	-34.74	-21.69	44.71	15.00

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152	ASP CO	-34.02	-21.34	43.43	15.00
152	ASP OF	-34.60	-21.55	42.34	15.00
152	ASP OF	2 -32.87	-20.86	43.49	15.00
152	ASP C	-34.87	-24.07	44.02	15.00
152	ASP O	-33.84	-24.43	44.56	15.00
153	GLU N	-35.33	-24.60	42.90	15.00
153	GLU CA	-34.70	-25.74	42.26	15.00
153	GLU CB	-35.49	-26.18	41.03	15.00
153	GLU CG	-35.79	-25.08	40.01	15.00
153	GLU CD	-37.17	-24.43	40.19	15.00
153	GLU OE	1 -38.12	-24.84	39.48	15.00
153	GLU OE	2 -37.29	-23.50	41.02	15.00
153	GLU C	-33.22	-25.56	41.91	15.00
153	GLU O	-32.59	-26.50	41.44	15.00
154	SER N	-32.66	-24.37	42.12	15.00
154	SER CA		-24.12	41.82	15.00
154	SER CB	-31.02	-22.70	41.29	15.00
154	SER OG	-32.05	-22.32	40.40	15.00
154	SER C	-30.40	-24.30	43.08	15.00
154	SER O	-29.17	-24.21	43.04	15.00
	CYS N	-31.08	-24.48	44.22	15.00
	CYS CA	-30.41	-24.66	45.49	15.00
	CYS C	-29.57	-25.91	45.40	15.00
	CYS O	-30.00	-26.91	44.82	15.00
	CYS CB	-31.44	-24.79	46.62	15.00
	CYS SG	-30.90	-24.05	48.19	15.00
	ASN N	-28.36	-25.87	45.94	15.00
	ASN CA	-27.47	-27.01	45.87	15.00
	ASN CB	-26.18	-26.59	45.19	15.00
	ASN CG	-25.28	-27.76	44.91	15.00
	ASN OD		-28.91	44.88	15.00
	ASN ND2		-27.49	44.73	15.00
	ASN C	-27.20	-27.67	47.23	15.00
	asn o	-26.43		48.05	15.00
	SER N	-27.79	-28.85	47.41	15.00
	SER CA	-27.67	-29.64	48.62	15.00
	SER CB	-28.48	-30.92	48.45	15.00
	SER OG	-29.83	-30.63	48.13	15.00
	SER C	-26.24	-30.00	49.04	15.00
	SER O	-26.01	-30.36	50.19	15.00
	ASP N		-29.93	48.10	15.00
	ASP CA	-23.89			15.00
158 .	ASP CB	-23.21	-30.69	47.07	15.00

158	ASP	CG	-23.72	-32.01	46.57	15.00
158	ASP	OD1	-24.95	-32.15	46.33	15.00
158	ASP	OD2	-22.86	-32.91	46.38	15.00
158	ASP	С	-23.14	-29.02	48.87	15.00
158	ASP	0	-22.11	-29.15	49.53	15.00
159	ASN	N	-23.62	-27.84	48.51	15.00
159	ASN	CA	-22.98	-26.62	48.94	15.00
159	ASN	CB	-23.10	-25.57	47.84	15.00
159	ASN	CG	-22.14	-24.40	48.03	15.00
159	ASN	OD1	-21.43	-24.30	49.04	15.00
159	ASN	ND2	-22.10	-23.51	47.04	15.00
159	ASN	С	-23.68	-26.14	50.20	15.00
159	ASN	0	-24.63	-25.37	50.12	15.00
160	LEU	N	-23.25	-26.64	51.36	15.00
160	LEU	CA	-23.84	-26.23	52.63	15.00
160	LEU	CB	-23.88	-27.42	53.60	15.00
160	LEU	CG	-24.59	-28.72	53.22	15.00
160	LEU	CD1	-24.45	-29.70	54.37	15.00
160	LEU	CD2	-26.06	-28.49	52.89	15.00
160	LEU	С	-22.98	-25.11	53.22	15.00
160	LEU	0	-21.91	-25.35	53.78	15.00
	asn		-23.47	-23.89	53.16	15.00
	ASN	CA	-22.70	-22.75	53.65	15.00
161		CB	-22.58	-21.73	52.53	15.00
161			-23.89	-21.51	51.84	15.00
	ASN		-24.74	-20.76	52.33	15.00
	ASN	ND2	-24.10	-22.23	50.75	15.00
	ASN		-23.23	-22.05	54.89	15.00
	ASN		-22.45	-21.62	55.73	15.00
	HIS		-24.54	-21.93	55.01	15.00
	HIS		-25.13	-21.27	56.16	15.00
162	HIS		-26.18	-20.27	55.67	15.00
	HIS		-26.55	-19.22	56.67	15.00
	HIS			-18.61	56.93	15.00
		ND1	-25.63		57.53	15.00
	HIS	CE1		-17.74	58.27	15.00
162	HIS	NE2		-17.69	57.93	15.00
		С	-25.76	-22.27	57.12	15.00
	HIS			-23.19	56.69	15.00
	ALA			-22.12	58.41	15.00
	ALA			-23.00	59.45	15.00
	ALA			-23.33	60.47	15.00
163	ALA	С	-27.15	-22.28	60.13	15.00

163 AI	A O	-27.00	-21.13	60.54	15.00
164 V	AL N	-28.30	-22.94	60.24	15.00
164 V	L CA	-29.48	-22.34	60.86	15.00
164 V	L CB	-30.54	-21.99	59.79	15.00
164 V	L CG1	-30.11	-20.79	58.99	15.00
164 V	L CG2	-30.75	-23.16	58.86	15.00
164 VA	ТС	-30.05	-23.28	61.91	15.00
164 VA	TO	-29.37	-24.21	62.33	15.00
165 LE	UN	-31.31	-23.07	62.30	15.00
165 LE	TU CA	-31.97	-23.88	63.33	15.00
165 LE	U CB	-32.00	-23.11	64.64	15.00
165 LE	TU CG	-32.59	-23.77	65.88	15.00
165 LE	U CD1	-31.53	-24.65	66.49	15.00
165 LE	U CD2	-33.04	-22.71	66.87	15.00
165 LE	UC	-33.40	-24.17	62.94	15.00
165 LE	UO	-34.16	-23.24	62.72	15.00
166 AL	A N	-33.79	-25.44	62.92	15.00
166 AL		-35.15	-25.82	62.56	15.00
166 AL		-35.16	-27.19	61.92	15.00
166 AL		-36.03	-25.80	63.80	15.00
	A O	-35.93	-26.66	64.66	15.00
167 VA		-36.89	-24.80	63.88	15.00
167 VA		-37.79	-24.62	65.01	15.00
167 VA		-38.11	-23.11	65.18	15.00
	L CG1	-39.35	-22.88	66.00	15.00
	L CG2	-36.94	-22.42	65.84	15.00
	L C	-39.06	-25.46	64.92	15.00
	r o	-39.83	-25.56	65.87	15.00
	YN	-39.28	-26.10	63.78	15.00
	Y CA	-40.48	-26.91	63.65	15.00
	YC	-40.77	-27.29	62.23	15.00
168 GL		-39.85	-27.36	61.41	15.00
169 TY		-42.04	-27.55	61.95	15.00
169 TY			-27.93	60.62	15.00
	R CB	-42.09	-29.38	60.30	15.00
	R CG	-42.61	-30.41	61.28	15.00
	R CD1	-43.94	-30.82	61.25	15.00
	R CE1	-44.42	-31.76	62.13	15.00
	R CD2	-41.77	-30.98	62.23	15.00
	R CE2	-42.24	-31.93	63.13	15.00
	R CZ	-43.57	-32.31	63.07	15.00
	R OH	-44.04	-33.26	63.95	15.00
169 TYI	( C	-44.02	-27.78	60.54	15.00

169 TYR O	-44.70	-27.74	61.57	15.00
170 GLY N	-44.55	-27.71	59.33	15.00
170 GLY CA	-45.98	-27.55	59.16	15.00
170 GLY C	-46.42	-27.49	57.72	15.00
170 GLY O	-45.69	-27.89	56.82	15.00
171 ILE N	-47.61	-26.95	57.49	15.00
171 ILE CA	-48.20	-26.83	56.16	15.00
171 ILE CB	-49.51	-27.69	56.06	15.00
171 ILE CG2	-50.16	-27.55	54.70	15.00
171 ILE CG1	-49.20	-29.18	56.27	15.00
171 ILE CD1	-48.97	-29.59	57.73	15.00
171 ILE C	-48.55	-25.36	55.90	15.00
171 ILE O	-48.58	-24.55	56.84	15.00
172 GLN N	-48.73	-25.00	54.63	15.00
172 GLN CA	-49.10	-23.64	54.25	15.00
172 GLN CB	-47.90	-22.90	53.68	15.00
172 GLN CG	-47.16	-22.04	54.69	15.00
172 GLN CD	-47.81	-20.68	54.87	15.00
172 GLN OE1	-47.24	-19.66	54.49	15.00
172 GLN NE2	-49.00	-20.66	55.47	15.00
172 GLN C	-50.24	-23.71	53.24	15.00
172 GLN 0	-51.31	-24.23	53.55	15.00
173 LYS N	-50.05	-23.17	52.05	15.00
173 LYS CA	-51.11	-23.24	51.05	15.00
173 LYS CB	-51.04	-22.05	50.08	15.00
173 LYS CG	-51.15	-20.70	50.77	15.00
173 LYS CD	-50.94	-19.57	49.77	15.00
173 LYS CE	-50.57	-18.27	50.50	15.00
173 LYS NZ	-49.28	-18.39	51.26	15.00
173 LYS C	-50.82	-24.55	50.34	15.00
173 LYS O	-50.33	-24.58	49.21	15.00
174 GLY N	-51.02	-25.63	51.08	15.00
174 GLY CA	-50.77	-26.96	50.56	15.00
174 GLY C	-49.30	-27.32	50.59	15.00
174 GLY 0			50.31	
175 ASN N		-26.35		15.00
175 ASN CA			50.98	
175 ASN CB		-25.39		
175 ASN CG		-24.86	49.14	15.00
175 ASN OD1		-23.79	49.08	15.00
175 ASN ND2			48.06	
175 ASN C			52.38	
175 ASN 0	-46.66	-26.16	53.33	15.00

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176 LYS N	-45.90	-28.13	52.53	15.00
176 LYS CA	-45.33	-28.50	53.82	15.00
176 LYS CB	-44.94	-29.98	53.84	15.00
176 LYS CG	-46.10	-30.95	53.63	15.00
176 LYS CD	-45.67	-32.36	53.98	15.00
176 LYS CE	-46.71	-33.40	53.61	15.00
176 LYS NZ	-46.36	-34.11	52.34	15.00
176 LYS C	-44.08	-27.63	53.94	15.00
176 LYS O	-43.52	-27.21	52.92	15.00
177 HIS N	-43.62	-27.37	55.16	15.00
177 HIS CA	-42.44	-26.52	55.34	15.00
177 HIS CB	-42.84	-25.05	55.23	15.00
177 HIS CG	-43.71	-24.59	56.35	15.00
177 HIS CD2	-43.41	-24.18	57.61	15.00
177 HIS ND1	-45.09	-24.52	56.26	15.00
177 HIS CE1	-45.59	-24.10	57.40	15.00
177 HIS NE2	-44.59	-23.89	58.24	15.00
177 HIS C	-41.74	-26.73	56.67	15.00
177 HIS O	-42.32	-27.24	57.62	15.00
178 TRP N	-40.49	-26.28	56.73	15.00
178 TRP CA	-39.64	-26.34	57.91	15.00
178 TRP CB	-38.24	-26.82	57.54	15.00
178 TRP CG	-38.16	-28.22	57.08	15.00
178 TRP CD2	-38.28	-29.41	57.88	15.00
178 TRP CE2	-38.08	-30.51	57.03	15.00
178 TRP CE3	-38.52	-29.64	59.24	15.00
178 TRP CD1	-37.92	-28.65	55.81	15.00
178 TRP NE1	-37.87	-30.02	55.77	15.00
178 TRP CZ2	-38.13	-31.83	57.48	15.00
178 TRP CZ3	-38.57	-30.95	59.70	15.00
178 TRP CH2	-38.37	-32.03	58.83	15.00
178 TRP C	-39.53	-24.91	58.39	15.00
178 TRP 0	-39.15	-24.04	57.61	15.00
179 ILE N	-39.89	-24.63	59.63	15.00
179 ILE CA	-39.79	-23.28	60.16	15.00
179 ILE CB	-40.65	-23.09	61.42	15.00
179 ILE CG2	-40.61	-21.64	61.85	15.00
179 ILE CG1	-42.09	-23.52	61.15	15.00
179 ILE CD1	-42.97	-23.45	62.38	15.00
179 ILE C	-38.32	-23.04	60.52	15.00
179 ILE O	-37.80	-23.66	61.45	15.00
180 ILE N	-37.67	-22.16	59.78	15.00
180 ILE CA	-36.27	-21.87	60.01	15.00

180	ILE	CB	-35.46	-22.00	58.70	15.00
180	ILE	CG2	-34.01	-21.64	58.91	15.00
180	ILE	CG1	-35.57	-23.42	58.16	15.00
180	ILE	CD1	-34.96	-24.46	59.08	15.00
180	ILE	С	-36.01	-20.52	60.69	15.00
180	ILE	0	-36.70	-19.53	60.44	15.00
181	LYS	N	-35.02	-20.53	61.58	15.00
181	LYS	CA	-34.60	-19.36	62.34	15.00
181	LYS	CB	-34.59	-19.69	63.84	15.00
181	LYS	CG	-34.02	-18.59	64.70	15.00
181	LYS	CD	-33.87	-19.04	66.14	15.00
181	LYS	CE	-33.58	-17.85	67.01	15.00
181	LYS	NZ	-33.52	-18.17	68.45	15.00
181	LYS	С	-33.19	-19.02	61.91	15.00
181	LYS	0	-32.28	-19.85	62.04	15.00
182	asn	N	-33.00	-17.82	61.37	15.00
182	ASN	CA	-31.68	-17.42	60.93	15.00
182	ASN	CB	-31.77	-16.80	59.54	15.00
182	ASN	CG	-30.45	-16.80	58.83	15.00
182	ASN	OD1	-29.40	-16.99	59.44	15.00
182	ASN	ND2	-30.48	-16.62	57.51	15.00
182	ASN	C	-31.10	-16.42	61.92	15.00
182	ASN	0	-31.81	-15.94	62.79	15.00
183	SER	N	-29.81	-16.14	61.82	15.00
183	SER	CA	-29.19	-15.20	62.74	15.00
183	SER	CB	-27.97	-15.85	63.38	15.00
183	SER	OG	-27.30	-16.66	62.44	15.00
183	SER	С	-28.79	-13.90	62.04	15.00
183	SER	0	-27.68	-13.40	62.23	15.00
184	TRP	N	-29.70	-13.34	61.25	15.00
184	TRP	CA	-29.44	-12.11	60.52	15.00
184	TRP	CB	-29.77	-12.29	59.03	15.00
184	TRP	CG	-28.79	-13.14	58.29	15.00
184	TRP	CD2	-28.97	-13.73	57.01	15.00
	TRP	CE2		-14.45	56.71	15.00
184				-13.74	56.08	15.00
184				-13.50	58.71	15.00
184	TRP	NE1		-14.28	57. <i>77</i>	15.00
184	TRP	CZ2	-27.64	-15.17	55.52	15.00
184	TRP		-29.85	-14.46	54.90	15.00
184				-15.16	54.63	15.00
1.84		С	-30.23	-10.93	61.07	15.00
184	TRP	0	-30.32	-5.88	60.43	15.00

			IADLL V.		
185 GLY	N -3	0.78 -	11.07	62.27	15.00
185 GLY	CA -3:	1.54	-9.98	62.83	15.00
185 GLY	C -3:	3.02 -	10.16	62.62	15.00
185 GLY		3.46 -	11.00	61.84	15.00
186 GLU		3.80	-9.35	63.33	15.00
186 GLU	CA -35	5.26	-9.39	63.27	15.00
186 GLU	CB -35	5.79	-8.71	64.53	15.00
186 GLU (	CG -37	7.29	-8.65	64.70	15.00
186 GLU (	CD -37	7.71	-8.03	66.04	15.00
186 GLU (	DE1 -36	.83	-7.61	66.83	15.00
186 GLU (	DE2 -38	.93	-7.98	66.31	15.00
186 GLU (	-35	.73	-8.65	62.03	15.00
186 GLU (	-36	.89	-8.73	61.64	15.00
187 ASN 1	v -34	.78	-8.02	61.36	15.00
187 ASN (	CA -35	.02	-7.21	60.18	15.00
187 ASN (	CB -34	.03	-6.03	60.25	15.00
187 ASN (	CG -34	.42	-4.86	59.37	15.00
187 ASN 0	DD1 -33	.58 -	-4.33	58.64	15.00
187 ASN N	TD2 -35	.67	-4.40	59.48	15.00
187 ASN C	-34	.86 -	-7.97	58.86	15.00
187 ASN C	-34	.92 -	-7.36	57.80	15.00
188 TRP N	ı –34	.62 -	-9.28	58.92	15.00
188 TRP C	A -34	.47 -1	10.08	57.70	15.00
188 TRP C	:B -33	.20 -1	LO.94	57.77	15.00
188 TRP C	:G -33	.05 -1	11.85	56.60	15.00
188 TRP C	D2 -33	.41 -1	13.23	56.52	15.00
	E2 -33	.18 -1	3.66	55.20	15.00
188 TRP C	E3 -33	.92 -1	4.16	57.45	15.00
188 TRP C	D1 -32	.61 -1	.1.50	55.36	15.00
188 TRP N	E1 -32	.69 -1	.2.58	54.51	15.00
188 TRP C			4.96	54.76	15.00
	<b>Z3</b> -34	.18 -1	.5.46	57.02	15.00
188 TRP C			.5.85	55.69	15.00
188 TRP C	-35	.66 -1	1.00	57.51	15.00
188 TRP 0	-36	.23 -1	1.48	58.49	15.00
189 GLY N	-36	.021	1.29	56.27	15.00
189 GLY C	A -37	.14 -1	2.17	56.00	15.00
189 GLY C	-38	.37 -1	1.86	56.84	15.00
189 GLY O	-38	.70 -1	0.69	57.08	15.00
190 ASN N	-39	.02 -1	2.91	57.32	15.00
190 ASN C	A -40	.22 -1	2.78	58.13	15.00
190 ASN C	B -41	.13 -1	4.00	57.93	15.00
190 ASN C	G -42	.58 -1	3.74	58.32	15.00

190	ASN	OD1	-42.87	-13.07	59.31	15.00
190	ASN	ND2	-43.50	-14.27	57.53	15.00
190	ASN	С	-39.86	-12.63	59.61	15.00
190	ASN	0	-39.81	-13.61	60.35	15.00
191	LYS	N	-39.55	-11.41	60.02	15.00
191	LYS	CA	-39.19	-11.14	61.41	15.00
191	LYS	CB	-40.43	-11.19	62.29	15.00
191	LYS	CG	-41.44	-10.10	61.96	15.00
191	LYS	CD	-40.92	-8.71	62.35	15.00
191	LYS	CE	-41.18	-7.63	61.27	15.00
191	LYS	NZ	-40.19	-7.69	60.13	15.00
191	LYS	С	-38.10	-12.07	61.94	15.00
191	LYS	0	-38.11	-12.46	63.11	15.00
192	GLY	N	-37.15	-12.41	61.07	15.00
192	GLY	CA	-36.04	-13.26	61.47	15.00
192	GLY	С	-36.15	-14.73	61.13	15.00
192	GLY	0	-35.19	-15.48	61.32	15.00
193	TYR	N	-37.30	-15.16	60.63	15.00
193	TYR	CA	-37.50	-16.56	60.28	15.00
193	TYR	CB	-38.69	-17.14	61.04	15.00
	TYR		-38.47	-17.29	62.51	15.00
193	TYR	CD1	-38.57	-16.19	63.36	15.00
193	TYR	CE1	-38.34	-16.32	64.72	15.00
193	TYR		-38.13	-18.51	63.06	15.00
193	TYR		-37.91	-18.65	64.42	15.00
193	TYR		-38.01	-17.55	65.24	15.00
193	TYR		-37.78	-17.71	66.59	15.00
193	TYR		-37.76	-16.73	58.80	15.00
193	TYR		-37.97	-15.76	58.08	15.00
194	ILE		-37.78	-17.99	58.37	15.00
194	ILE		-38.03	-18.32	56.99	15.00
194	ILE		-36.75	-18.27	56.13	15.00
194	ILE		-35.65	-19.10	56.76	15.00
	ILE		-37.06			15.00
			-35.95		53.75	15.00
	ILE		-38.65		56.88	15.00 15.00
194	ILE		-38.20		57.51	15.00
195			-39.71	<b>-19.79</b>	56.09	
	LEU			-21.04 -20.78	55.87 55.71	15.00 15.00
	LEU			-20.78	55.71	15.00
	LEU			-20.89 -20.31	56.98 58.19	15.00
	LEU					15.00
エスコ	LEU	CD2	-44.09	-20.13	56.75	13.00

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195	LEU	С	-39.83	-21.68	54.62	15.00
195	LEU	0	-40.04	-21.19	53.51	15.00
196	MET	N	-39.05	-22.73	54.81	15.00
196	MET	CA	-38.44	-23.44	53.70	15.00
196	MET	CB	-37.01	-23.80	54.04	15.00
196	MET	CG	-36.11	-22.58	54.16	15.00
196	MET	SD	-34.46	-22.98	54.73	15.00
196	MET	CE	-33.78	-23.75	53.28	15.00
196	MET	С	-39.27	-24.68	53.38	15.00
196	MET	0	-40.03	-25.14	54.23	15.00
197	ALA	N	-39.15	-25.19	52.16	15.00
197	ALA	CA	-39.91	-26.36	51.72	15.00
197	ALA	CB	-39.86	-26.48	50.19	15.00
197	ALA	С	-39.51	-27.69	52.36	15.00
197	ALA	0	-38.33	-28.00	52.50	15.00
198	ARG	N	-40.52	-28.50	52.67	15.00
198	ARG	CA	-40.32	-29.81	53.28	15.00
198	ARG	CB	-41.08	-29.90	54.60	15.00
198	ARG	CG	-41.09	-31.28	55.23	15.00
	ARG		-41.40	-31.21	56.71	15.00
198	ARG	NE	-42.71	-30.64	57.00	15.00
198	ARG	CZ	-43.83	-31.35	57.04	15.00
198	ARG	NH1	-43.80	-32.66	56.80	15.00
	ARG		-44.97	-30.76	57.34	15.00
198	ARG		-40.79	-30.90	52.34	15.00
198	ARG	0	-41.87	-30.81	51.76	15.00
199	ASN		-39.97	-31.93	52.20	15.00
199	ASN		-40.28	-33.07	51.33	15.00
199	ASN		-41.68	-33.62	51.60	15.00
199	ASN		-41.76	-34.41	52.90	15.00
199	ASN		-42.80	-34.44	53.57	15.00
199	ASN		-40.65	-35.05	53.28	15.00
	ASN	С	-40.08	-32.78	49.85	15.00
		0		-33.59	48.99	15.00
	LYS	N	-39.47	-31.65	49.54	15.00
200	LYS		-39.18	-31.30	48.16	15.00
		CB	-39.18	-29.77	47.95	15.00
	LYS	CG	-40.55	-29.15	47.76	15.00
200		CD	-41.24	-29.74	46.54	15.00
200	LYS		-42.64	-29.19	46.34	15.00
200		NZ	-43.39	-29.98	45.32	15.00
	LYS		-37.80	-31.87	47.84	15.00
200	LYS	0	-36.86	-31.13	47.56	15.00

201	ASN	N	-37.66	-33.19	48.01	15.00
201	ASN	CA	-36.40	-33.87	47.72	15.00
201	ASN	CB	-36.24	-33.99	46.19	15.00
201	ASN	CG	-35.16	-35.00	45.77	15.00
201	ASN	OD1	-35.27	-35.62	44.69	15.00
201	ASN	ND2	-34.11	-35.15	46.58	15.00
201	ASN	С	-35.18	-33.16	48.34	15.00
201	ASN	0	-34.28	-32.70	47.62	15.00
202	ASN	N	-35.16	-33.09	49.67	15.00
202	ASN	CA	-34.05	-32.47	50.41	15.00
202	ASN	CB	-32.86	-33.43	50.47	15.00
202	ASN	CG	-31.99	-33.22	51.69	15.00
202	ASN	OD1	-32.43	-32.60	52.67	15.00
202	ASN	ND2	-30.77	-33.72	51.66	15.00
202	ASN	С	-33.61	-31.11	49.86	15.00
202	ASN	0	-32.42	-30.87	49.62	15.00
203	ALA	N	-34.58	-30.22	49.72	15.00
203	ALA	н	-35.43	-30.48	50.10	15.00
203	ALA	CA	-34.36	-28.88	49.17	15.00
203	ALA	CB	-35.62	-28.06	49.20	15.00
203	ALA	С	-33.31	-28.14	50.00	15.00
203	ALA	0	-33.47	-27.92	51.19	15.00
204	CYS	N	-32.23	-27.75	49.33	15.00
204	CYS	CA	-31.15	-27.01	49.98	15.00
204	CYS	С	-30.39	-27.80	51.03	15.00
204	CYS	0	-29.68	-27.21	51.83	15.00
204	CYS	CB	-31.68	-25.71	50.61	15.00
204	CYS	SG	-32.40	-24.49	49.47	15.00
205	GLY	N	-30.51	-29.13	51.01	15.00
205	GLY	CA	-29.80	-29.96	51.97	15.00
205	GLY	С	-30.19	-29.78	53.44	15.00
205	GLY	0	-29.39	-30.02	54.34	15.00
206	ILE	N	-31.43	-29.40	53.67	15.00
206	ILE	CA	-31.94	-29.18	55.01	15.00
206	ILE	CB	-33.46	-28.88	54.98	15.00
206	ILE	CG2	-34.18	-29.93	54.15	15.00
206	ILE	CG1	-34.03	-28.79	56.39	15.00
206	ILE	CD1	-33.55	-27.60	57.17	15.00
206	ILE	С	-31.63	-30.32	55.98	15.00
206	ILE	0	-31.31	-30.07	57.15	15.00
207	ALA	N	-31.68	-31.56	55.50	15.00
207	ALA	CA	-31.40	-32.72	56.36	15.00
207	ALA	CB	-32.50	-33.74	56.24	15.00

207	ALA C	-30.07	-33.37	56.02	15.00
207	7 ALA O	-29.89	-34.56	56.21	15.00
208	B ASN N	-29.11	-32.58	55.55	15.00
208	B ASN CA	-27.81	-33.10	55.16	15.00
208	ASN CB	-27.34	-32.39	53.88	15.00
208	ASN CG	-27.15	-33.35	52.72	15.00
208	ASN OD1	-28.12	~33.85	52.15	15.00
208	ASN ND2	-25.90	-33.60	52.36	15.00
208	ASN C	-26.72	-33.00	56.22	15.00
208	ASN O	-25.66	-33.62	56.10	15.00
209	LEU N	-26.96		57.27	15.00
209	LEU CA	-25.96	-32.06	58.31	15.00
209	LEU CB	-24.98	-30.96	57.89	15.00
209	LEU CG	-23.69	-30.69	58.67	15.00
209		-22.77	-31.90	58.61	15.00
	LEU CD2	-22.99		58.09	15.00
209		-26.63	· -	59.63	15.00
	LEU O	-26.11	-30.91	60.41	15.00
	ALA N	-27.79	-32.31	59.90	15.00
	ALA H	-28.15	-32.83	59.17	15.00
	ALA CA	-28.56	-32.01	61.10	15.00
210		-29.98		60.98	15.00
	ALA C	-27.93	-32.71	62.32	15.00
	ALA O	-27.23	-33.70	62.23	15.00
	SER N	-28.20	-32.12	63.50	15.00
211		-27.73	-32.66	64.78	15.00
211		-26.21	-32.55	64.89	15.00
211		-25.79	-31.20	64.98	15.00
211		-28.38	-31.89	65.92	15.00
211		-28.93	-30.80	65.72	15.00
212	PHE N	-28.35	-32.47	67.11	15.00
212	PHE CA	-28.93			15.00
212		-30.43	-32.16	68.36	15.00
	PHE CG		-33.63	68.34	15.00
	PHE CD1		-34.23	67.20	15.00
	PHE CD2	-30.55		69.48	15.00
212		-31.61	-35.57	67.19	15.00
212		-30.87	-35.79	69.48	15.00
212		-31.40	-36.35	68.33	15.00
212	PHE C		-32.36	69.50	15.00
212	PHE O		-33.48	69.48	15.00
213			-31.54	70.55	15.00
213	PRO CD	-28.53	-30.16	70.68	15.00

213	PRO	CA	-27.32	-31.95	71.76	15.00
213	PRO	CB	-26.95	-30.61	72.38	15.00
213	PRO	CG	-28.16	-29.81	72.11	15.00
213	PRO	С	-28.20	-32.77	72.70	15.00
213	PRO	0	-29.42	-32.64	72.69	15.00
214	LYS	N	-27.58	-33.60	73.53	15.00
214	LYS	CA	-28.32	-34.41	74.49	15.00
214	LYS	СВ	-27.85	-35.85	74.47	15.00
214	LYS	CG	-28.28	-36.60	73.23	15.00
214	LYS	CD	-27.98	-38.09	73.30	15.00
214	LYS	CE	-26.48	-38.39	73.31	15.00
214	LYS	NZ	-25.86	-38.24	74.66	15.00
214	LYS	С	-28.17	-33.84	75.89	15.00
214	LYS	0	-27.07	-33.47	76.29	15.00
215	MET	N	-29.28	-33.75	76.61	15.00
215	MET	CA	-29.29	-33.24	77.98	15.00
215	MET	CB	-30.27	-32.08	78.11	15.00
215	MET	CG	-29.79	-30.79	77.48	15.00
215	MET	SD	-28.97	-29.73	78.67	15.00
215	MET	CE	-30.38	-28.92	79.43	15.00
215	MET	С	-29.67	-34.33	78.99	15.00
215	MET	OT1	-30.25	-35.37	78.59	15.00
215	MET	OT2	-29.39	-34.13	80.20	15.00
216	HOH	OH2	-21.96	-40.63	81.12	15.00
217	HOH	OH2	-30.77	-17.16	67.86	15.00
218	HOH	OH2	-30.16	-20.07	64.02	15.00
219	HOH	OH2	-3.64	-10.82	59.75	15.00
220	HOH	OH2	-13.18	-7.77	71.57	15.00
221	HOH	OH2	-34.51	-22.61	70.17	15.00
222	HOH	OH2	-18.02	-34.44	65.29	15.00
223	HOH	OH2	-17.01	-5.28	69.42	15.00
224	HOH	OHS	-24.38	-30.77	62.26	15.00
225	HOH	OH2	0.36	-5.40	64.98	15.00
226	HOH	OH2	-13.68	-21.42	66.86	15.00
	HOH	OH2	-46.72	-29.80	50.41	15.00
228	HOH		-45.10	-36.23	56.40	15.00
229	HOH			-12.35	65.48	15.00
230	HOH		-35.85	-37.05	52.41	15.00
231	HOH	OH2	-19.20	-39.14	66.78	15.00
232	нон		-30.09	-19.72	66.64	15.00
233	HOH		-27.95	-19.50	62.38	15.00
234			-21.75	-30.29	62.28	15.00
235	HOH	OH2	-30.30	-2.55	77.57	15.00

9

236	нон	OH2	-33.08	-28.99	86.45	15.00
237	HOH	OH2	-30.07	-22.68	84.37	15.00
238	HOH	OH2	-39.83	-16.82	48.34	15.00
239	HOH	OH2	-34.57	-24.95	47.01	15.00
240	HOH	OH2	-46.44	-34.07	57.12	15.00
241	HOH	OH2	-26.91	-7.02	56.22	15.00
242	HOH	OH2	-42.10	-15.05	61.98	15.00
243	нон	OH2	-24.27	-7.11	65.05	15.00
244	нон	OH2	-33.44	-27.69	70.80	15.00
245	нон	OH2	-40.50	-27.38	80.61	15.00
246	нон	OH2	-14.45	-17.44	86.64	15.00
247	нон	OH2	-4.86	-12.23	73.56	15.00
248	нон	OH2	-10.86	-20.50	79.87	15.00
249	нон	OH2	-27.43	-35.04	59.25	15.00
250	нон	OH2	-35.26	-10.90	53.73	15.00
251	нон	OH2	-31.84	-29.20	46.92	15.00
252	HOH	OH2	-42.75	-9.71	40.49	15.00
253	HOH	OH2	-41.27	-34.56	56.25	15.00
254	нон	OH2	-44.55	-15.65	65.22	15.00
255	HOH	OH2	-32.52	-13.49	60.73	15.00
256	нон	OH2	-39.73	-4.62	63.34	15.00
257	нон	OH2	-25.69	-11.84	70.98	15.00
258	HOH	OH2	-31.93	-6.64	63.98	15.00
259	HOH	OH2	-19.62	-7.72	62.94	15.00
260	HOH	OH2	-33.42	-20.20	70.53	15.00
261	HOH	OH2	-12.62	-24.00	79.04	15.00
262	HOH	OH2	-9.78	-21.46	77.40	15.00
263	HOH	OH2	-6.71	-27.36	80.84	15.00
264	HOH	OH2	-21.06	-35.71	57.19	15.00
265	HOH	OH2	-26.47	-48.97	59.68	15.00
266	HOH	OH2	-14.22	-32.57	69.97	15.00
267	HOH	OH2	-11.69	-25.57	76.63	15.00
268	HOH	OH2	-17.38	-27.79	86.86	15.00
269	HOH	OH2	-22.39	-37.94	70.91	15.00
270	HOH	OH2	-10.44	-11.32	63.69	15.00
271	HOH	OH2	-8.66	-22.33	72.95	15.00
272	нон	OH2	-29.93	-20.17	48.73	15.00
273	HOH	OH2	-22.92	-30.27	39.30	15.00
274	HOH	OH2	-33.19	-37.20	49.46	15.00
275	нон	OH2	-28.10	-25.82	41.06	15.00
276	нон	OH2	-35.93	-29.91	44.54	15.00
277	нон	OH2	-37.76	-30.41	51.24	15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors  $({\tt A}^2)$  for the cathepsin K complex with inhibitor 4-[N-

Residue	Atom	x	Y	Z	В
1 ALA	СВ	-46.25	-39.17	62.96	30.60
1 ALA	C	-47.93	-37.51	63.80	29.74
1 ALA	0	-49.14	-37.57	63.58	32.13
1 ALA	N	-48.18	-39.83	64.36	28.23
1 ALA	CA	-47.15	-38.78	64.13	28.86
2 PRO	N	-47.26	-36.34	63.80	27.19
2 PRO	CD	-45.94	-36.10	64.40	26.45
2 PRO	CA	-47.92	-35.06	63.50	27.01
2 PRO	CB	-47.28	-34.10	64.52	26.65
· 2 PRO	CG	-46.25	-34.95	65.31	27.69
2 PRO	С	-47.73	-34.52	62.09	26.37
2 PRO	0	-46.67	-34.70	61.50	26.53
3 ASP	N	-48.76	-33.86	61.58	26.63
3 ASP	CA	-48.73	-33.23	60.26	24.49
3 ASP	CB	-50.14	-33.03	59.69	23.94
3 ASP	CG	-50.75	-34.32	59.17	24.73
3 ASP	OD1	-50.19	-34.88	58.21	31.10
3 ASP		-51.79	-34.76	59.71	23.79
3 ASP	С	-48.03	-31.88	60.39	24.62
3 ASP	0	-47.08	-31.59	59.67	23.92
4 SER	N	-48.55	-31.04	61.28	24.05
4 SER	CA	-47.98	-29.72	61.55	22.83
4 SER	CB	-49.04	-28.62	61.52	23.29
4 SER	OG	-49.94	-28.70	60.36	24.54
4 SER	C	-47.30	-29.75	62.91	23.31
4 SER	0	-47.71	-30.51	63.79	26.63
5 VAL	N	-46.27	-28.92	63.09	24.43
5 VAL	CA	-45.52	-28.80	64.34	22.41
5 VAL		-44.44	-29.91	64.50	24.60
5 VAL	CG1	-43.39	-29.50	65.53	19.58
5 VAL		-45.09	-31.22	64.94	26.54
5 VAL	C	-44.80	-27.45	64.30	22.72

5	VAL	0	-44.23	-27.09	63.27	24.78
6	ASP	N	-44.80	-26.74	65.41	22.25
6	ASP	CA	-44.17	-25.43	65.48	20.13
6	ASP	CB	-45.15	-24.34	65.04	20.94
6	ASP	CG	-44.49	-22.99	64.81	19.46
6	ASP	OD1 ·	-43.28	-22.84	65.12	15.02
6	ASP	OD2	-45.20	-22.09	64.31	15.42
6	ASP	С	-43.67	-25.17	66.88	20.15
6	ASP	0	-44.45	-24.78	67.74	26.96
7	TYR	N	-42.37	-25.32	67.10	20.70
7	TYR	CA	-41.82	-25.08	68.42	19.50
7	TYR	CB	-40.44	-25.71	68.58	22.46
7	TYR	CG	-40.51	-27.21	68.76	22.69
7	TYR	CD1	-40.50	-27.78	70.03	24.39
7	TYR	CE1	-40.61	-29.16	70.20	23.01
7	TYR	CD2	-40.62	-28.05	67.66	17.84
7	TYR	CE2	-40.72	-29.42	67.82	17.55
7	TYR	CZ	-40.72	-29.97	69.08	16.70
7	TYR	OH	-40.84	-31.33	69.22	21.92
7	TYR	С	-41.77	-23.62	68.78	18.59
7	TYR	0	-41.12	-23.25	69.75	23.10
8	ARG	N	-42.39	-22.78	67.96	16.73
8	ARG	CA	-42.44	-21.36	68.25	17.40
8	ARG	CB	-42.56	-20.52	66.98	19.28
8	ARG	CG	-41.27	-20.40	66.17	22.66
8	ARG	CD	-41.41	-19.48	64.98	14.98
8	ARG	NE	-42.40	-19.99	64.04	17.92
8	ARG	CZ	-42.59	-19.53	62.81	22.05
8	ARG	NHl	-41.85	-18.52	62.36	24.53
8	ARG	NH2	-43.50	-20.10	62.03	26.01
8	ARG	С	-43.64	-21.14	69.17	15.35
8	ARG	0	-43.52	-20.55	70.24	14.09
9	LYS	N	-44.77	-21.73	68.79	13.90
9	LYS	CA	-46.01	-21.63	69.54	13.24
9	LYS	CB	-47.21	-21.84	68.62	10.93
9	LYS	CG	-47.27	-20.83	67.48	16.03
9	LYS	CD	-48.31	-21.19	66.44	13.55
9	LYS	CE	-49.69	-21.20	67.04	17.52
9	LYS	NZ	-50.71	-21.24	65.97	18.86
9	LYS	С	-46.01	-22.61	70.70	15.37
9	LYS	0	-47.06	-23.10	71.14	15.03
0	LYS	N	-44.81	-22.90	71.19	17.15

10	LYS	CA	-44.58	-23.82	72.31	14.97
10	LYS	CB	-44.10	-25.19	71.81	13.68
10	LYS	CG	-45.15	-26.06	71.14	17.87
10	LYS	CD	-44.58	-27.44	70.87	17.96
10	LYS	CE	-45.67	-28.49	70.71	22.08
10	LYS	NZ	-45.07	-29.82	70.35	25.39
10	LYS	С	-43.52	-23.20	73.23	10.83
10	LYS	0	-43.08	-23.82	74.19	9.39
11	GLY	N	-43.07	-22.01	72.88	11.15
11	GLY	CA	-42.08	-21.35	73.70	14.48
11	GLY	С	-40.71	-21.99	73.69	15.75
11	GLY	0	-39.92	-21.75	74.60	16.24
12	TYR	N	-40.41	-22.80	72.67	16.05
12	TYR	CA	-39.09	-23.44	72.59	15.26
12	TYR	CB	-39.18	-24.83	71.96	16.85
12	TYR	CG	-39.84	-25.90	72.80	12.79
12	TYR	CD1	-41.22	-26.04	72.82	12.72
12	TYR	CE1	-41.82	-27.07	73.52	16.82
12	TYR	CD2	-39.08	-26.82	73.50	18.15
12	TYR	CE2	-39.67	-27.86	74.20	21.62
12	TYR	CZ	-41.04	-27.98	74.20	19.19
12	TYR	OH	-41.63	-29.03	74.89	22.22
12	TYR	С	-38.08	-22.61	71.81	16.45
12	TYR	0	-36.89	-22.92	71.82	18.97
13	VAL	N	-38.55	-21.55	71.16	16.26
13	VAL	CA	-37.67	-20.71	70.34	17.11
13	VAL	CB	-38.25	-20.58	68.90	17.86
13	VAL	CG1	-37.20	-20.07	67.96	18.97
13	VAL	CG2	-38.77	-21.92	68.41	19.17
13	VAL	C	-37.37	-19.31	70.89	16.02
13	VAL	0	-38.27	-18.51	71.14	16.61
14	THR	N	-36.08	-19.01	71.03	15.90
14	THR	CA	-35.65	-17.71	71.53	20.92
14	THR	CB	-34.21	-17.77	72.04	22.63
14	THR	OG1	-33.33	-18.11	70.97	28.59
14	THR	CG2	-34.09	-18.78	73.17	23.81
14	THR	C	-35.73	-16.67	70.41	25.69
14	THR	0	-36.01	-17.02	69.26	28.47
15	PRO	N	-35.50	-15.37	70.73	28.26
15	PRO	CD	-35.15	-14.82	72.05	27.19
15	PRO	CA		-14.29	69.73	23.60
15	PRO	CB	-35.29	-13.04	70.58	24.84

15 PRO	CG	-34.43	-13.55	71.68	26.75
15 PRO	С	-34.53	-14.44	68.62	19.41
15 PRO	0	-33.43	-14.94	68.84	15.67
16 VAL	N	-34.90	-13.96	67.43	16.42
16 VAL	CA	-34.03	-14.06	66.27	12.13
16 VAL	CB	-34.77	-13.71	64.95	9.20
16 VAL	CG1	-33.84	-13.89	63.77	7.84
16 VAL	CG2	-36.01	-14.56	64.78	6.59
16 VAL	С	-32.81	-13.16	66.39	11.22
16 VAL	0	-32.93	-11.94	66.44	11.47
17 LYS	N	-31.65	-13.79	66.45	10.35
17 LYS	CA	-30.39	-13.07	66.55	9.97
17 LYS	CB	-29.31	-13.93	67.22	14.75
17 LYS	CG	-29.17	-13.71	68.75	10.18
17 LYS	CD	-30.45	-14.02	69.52	5.48
17 LYS	CE	-30.69	-15.51	69.66	15.18
17 LYS	NZ	-29.75	-16.23	70.59	13.03
17 LYS	С	-29.98	-12.57	65.16	9.61
17 LYS	0	-30.72	-12.73	64.20	9.93
18 ASN 1	N	-28.76	-12.05	65.05	8.91
18 ASN (	CA	-28.28	-11.47	63.80	8.23
18 ASN (	CB	-28.78	-10.03	63.72	10.88
18 ASN (	CG	-28.51	-9.36	62.39	13.91
18 ASN (	OD1	-27.51	-9.63	61.72	9.48
18 ASN 1	ND2	-29.40	-8.44	62.03	13.48
18 ASN (	С	-26.75	-11.54	63.75	8.02
18 ASN (		-26.07	-10.64	64.22	13.56
19 GLN 1		-26.22	-12.59	63.14	7.59
19 GLN (	CA	-24.78	-12.81	63.06	9.07
19 GLN (		-24.50	-14.08	62.26	11.76
19 GLN (		-24.91	-14.00	60.81	10.58
19 GLN (		-24.73	-15.33	60.11	10.05
19 GLN (		-25.69	-16.07	59.88	8.08
19 GLN 1			-15.66	59.80	3.31
19 GLN (		-23.84		62.60	12.84
19 GLN (		-22.65	-11.71	62.95	14.67
20 GLY 1		-24.33	-10.75	61.81	9.97
20 GLY (		-23.45	-9.68	61.35	9.75
20 GLY (		-22.40	-10.16	60.37	9.12
		-22.56	-11.21	59.74	14.46
21 GLN 1			-9.40	60.20	10.42
21 GLN (	CA	-20.28	-9.80	59.25	13.95

21	GLN	CB	-19.55	-8.58	58.66	13.14
21	GLN	CG	-20.40	-7.79	57.65	12.44
21	GLN	CD	-20.73	-8.61	56.41	13.48
21	GLN	OE1	-19.84	-9.15	55.76	14.90
21	GLN	NE2	-22.02	-8.71	56.08	9.41
21	GLN	С	-19.30	-10.83	59.81	15.26
21	GLN	0	-18.08	-10.64	59.79	16.33
22	CYS	N	-19.86	-11.93	60.29	17.72
22	CYS	CA	-19.10	-13.04	60.86	16.36
22	CYS	С	-19.82	-14.31	60.40	16.07
22	CYS	0	-21.05	-14.39	60.44	8.26
22	CYS	CB	-19.02	-12.91	62.40	16.53
22	CYS	SG	-18.36	-14.33	63.35	15.48
23	GLY	N	-19.04	-15.25	59.83	16.86
23	GLY	CA	-19.59	-16.52	59.35	14.89
23	GLY	С	-19.67	-17.49	60.52	13.50
23	GLY	0	-18.91	-18.45	60.61	11.64
24	SER	N	-20.61	-17.20	61.41	13.66
24	SER	CA	-20.82	-17.99	62.61	14.12
24	SER	CB	-20.65	-17.10	63.84	17.06
24	SER	OG	-21.37	-15.88	63.67	19.58
24	SER	С	-22.18	-18.67	62.64	14.42
24	SER	0	-22.63	-19.12	63.69	15.12
25	CYS	N	-22.83	-18.77	61.48	15.74
25	CYS	CA	-24.16	-19.38	61.40	12.45
25	CYS	CB	-24.61	-19.48	59.92	17.82
25	CYS	SG	-23.46	-20.34	58.77	15.84
25	CYS	C	-24.23	-20.73	62.12	12.21
25	CYS	0	-25.27	-21.09	62.66	8.88
25	INH	C1	-26.76	-10.18	57.23	37.63
25	INH	C2	-25.50	-10.64	57.58	37.16
25	INH	C3	-24.85	-11.61	56.79	34.05
25	INH		-25.45	-12.12	55.64	32.87
25	INH	C5	-26.72	-11.65	55.30	36.05
25.	INH	C6	-27.38	-10.68	56.09	37.28
25	INH	C7	-24.76	-13.16	54.79	31.70
25	INH	08	-24.07	-14.24	55.46	33.18
25	INH	C9	-24.20	-15.65	55.36	32.90
25	INH	010	-24.83	-16.33	56.19	27.65
25	INH	C11	-23.57	-17.64	54.11	33.43
25	INH	C12	-23.56	-17.98	52.63	29.93
25	INH	C13	-24.79	-17.58	51.82	30.09

25	INH	C14	-24.84	-16.08	51.57	28.76
25	INH	C15	-24.70	-18.31	50.53	33.84
25	INH	C16	-22.36	-18.24	54.80	34.35
25	INH	017	-21.26	-18.25	54.27	39.78
25	INH	N18	-22.58	-18.72	56.02	35.16
25	INH	C19	-21.64	-19.29	56.85	29.32
25	INH	N20	-23.57	-16.20	54.33	34.33
25	INH	C21	-21.16	-20.68	56.54	29.68
25	INH	C22	-22.10	-19.32	58.29	26.25
25	INH	023	-22.30	-18.20	58.72	26.33
25	INH	C24	-13.39	-26.50	60.06	25.12
25	INH	C25	-13.12	-25.19	59.68	26.39
25	INH	C26	-14.04	-24.48	58.91	25.10
25	INH	C27	-15.23	-25.07	58.52	23.88
25	INH	C28	-15.49	-26.37	58.90	24.81
25	INH	C29	-14.58	-27.09	59.67	23.22
25	INH	C30	-16.20	-24.31	57.66	25.64
25	INH	031	-16.99	-24.94	56.63	26.66
25	INH	C32	-18.41	-24.84	56.56	24.33
25	INH		-19.08	-25.66	55.96	24.89
25	INH		-20.39	-23.56	57.26	25.72
25	INH	C35	-21.15	-24.78	57.78	22.12
25	INH	C36	-21.80	-25.75	56.80	16.61
25	INH		-23.15	-25.25	56.41	15.08
25	INH		-21.91	-27.08	57.47	16.77
25	INH		-20.56	-22.41	58.25	28.82
25	INH		-20.39	-22.62	59.45	37.25
25	INH	N41	-20.88	-21.18	57.82	28.59
25	INH		-21.00	-20.17	58.81	27.26
25	INH		-18.95	-23.81	57.20	25.99
26	TRP		-23.11	-21.45	62.14	14.10
26	TRP		-23.02	-22.74	62.82	15.29
26	TRP		-21.66	-23.38	62.56	12.22
	TRP		-20.53			15.74
			-19.74		64.10	17.65
		CE2	-18.76		63.98	15.49
		CE3	-19.77	-23.30	65.27	18.53
	TRP		-20.02	-21.49	62.14	16.57
	TRP		-18.95	-20.90	62.77	17.92
	TRP		-17.82		64.98	13.72
26				-23.05		16.53
26	TRP	CH2	-17.87	-22.04	66.11	15.09

26	TRP	С	-23.24	-22.55	64.32	19.46
26	TRP	0	-24.09	-23.21	64.92	24.77
27	ALA	N	-22.52	-21.59	64.90	19.49
27	ALA	CA	-22.61	-21.25	66.31	12.94
27	ALA	CB	-21.77	-20.01	66.61	12.05
27	ALA	С	-24.07	-20.99	66.64	8.35
27	ALA	0	-24.61	-21.54	67.60	6.75
28	PHE	N	-24.72	-20.18	65.80	8.07
28	PHE	CA	-26.13	-19.83	65.95	9.51
28	PHE	CB	-26.51	-18.67	65.04	7.62
28	PHE	CG	-25.96	-17.35	65.48	4.72
28	PHE	CD1	-24.74	-16.91	65.01	4.11
28	PHE	CD2	-26.66	-16.56	66.38	2.92
28	PHE	CE1	-24.22	-15.69	65.41	4.53
28	PHE	CE2	-26.16	-15.33	66.79	2.38
28	PHE	CZ	-24.93	-14.89	66.31	2.00
28	PHE	С	-27.07	-21.01	65.72	10.66
28	PHE	0	-28.18	-21.04	66.26	14.07
29	SER	N	-26.64	-21.96	64.89	10.83
29	SER	CA	-27.44	-23.15	64.62	8.54
29	SER	CB	-26.92	-23.88	63.37	2.45
29	SER	OG	-27.80	-24.93	62.97	2.00
29	SER	C	-27.40	-24.05	65.86	7.86
29	SER	0	-28.44	-24.46	66.38	6.37
30	SER	N	-26.19	-24.29	66.36	5.14
30	SER	CA	-26.00	-25.14	67.52	7.65
30	SER	CB	-24.52	-25.27	67.84	10.56
30	SER	OG	-23.81	-25.61	66.67	14.28
30	SER	C	-26.76	-24.61	68.73	6.72
30	SER	0	-27.57	-25.32	69.34	8.64
31	VAL	N	-26.50	-23.35	69.06	6.44
31	VAL	CA	-27.15	-22.71	70.19	6.61
31	VAL	CB	-26.73	-21.23	70.25	6.76
31	VAL	CG1		-20.40	71.03	9.71
31	VAL	CG2	-25.35	-21.14	70.90	2.00
31	VAL	С	-28.67	-22.86	70.18	9.86
31	VAL	0	-29.25	-23.30	71.17	13.64
32	GLY	N	-29.30	-22.56	69.05	12.98
32	GLY	CA	-30.75	-22.68	68.94	9.41
32	GLY	С	-31.27	-24.10	69.19	10.59
32	GLY	0		-24.29	69.59	8.20
33	ALA	N	-30.44	-25.10	68.91	11.42

33	ALA	CA	-30.82	-26.50	69.12	13.15
33	ALA	CB	-29.82	-27.42	68.47	10.51
33			-30.86	-26.73	70.64	14.58
33	ALA	0	-31.87	-27.19	71.19	12.75
34	LEU	I N	-29.75	-26.39	71.29	12.77
34	LEU	CA	-29.62	-26.51	72.73	12.38
34	LEU	CB	-28.26	-26.01	73.20	4.32
34	LEU	CG	-27.04	-26.64	72.57	4.80
34	LEU	CD1	-25.82	-25.86	72.95	6.27
34	LEU	CD2	-26.92	-28.09	73.01	4.63
34	LEU	C	-30.73	-25.73	73.43	16.38
34	LEU	0	-31.32	-26.22	74.39	18.32
35	GLU	N	-31.00	-24.53	72.95	15.34
35	GLU	CA	-32.03	-23.71	73.54	15.20
35	GLU	CB	-32.17	-22.38	72.81	12.21
35	GLU	CG	-30.92	-21.55	72.88	18.17
35	GLU	CD	-31.07	-20.22	72.17	21.50
35	GLU	OE1	-31.87	-20.14	71.21	23.90
35	GLU	OE2	-30.40	-19.25	72.57	22.09
35	GLU	С	-33.37	-24.43	73.60	18.81
35	GLU	0	-34.06	-24.37	74.62	23.06
36	GLY	N	-33.73	-25.11	72.52	19.33
36	GLY	CA	-35.00	-25.83	72.48	19.77
36	GLY	С	-34.96	-27.02	73.42	21.74
36	GLY	0	-35.97	-27.38	74.05	18.02
37	GLN	N	-33.79	-27.63	73.53	23.70
37	GLN	CA	-33.61	-28.77	74.41	23.25
37	GLN	CB	-32.27	-29.48	74.15	22.37
37	GLN	CG	-32.08	-29.98	72.71	26.26
37	GLN	CD	-33.38	-30.28	71.94	30.06
37	GLN	OE1	-34.20	-31.11	72.36	29.64
37	GLN	NE2	-33.58	-29.57	70.83	31.26
37	GLN		-33.73	-28.32	75.86	24.67
37	GLN	0	-34.51	-28.89	76.62	25.08
38	LEU	N	-32.99	-27.27	76.22	22.29
38	LEU	CA	-33.04	-26.73	77.57	23.08
38	LEU	CB	-32.20	-25.46	77.70	21.38
38	LEU	CG	-32.11	-24.78	79.07	17.48
38	LEU	CD1	-31.77	-25.77	80.17	13.71
38	LEU	CD2	-31.07	-23.68	78.99	17.90
38	LEU	С	-34.47	-26.46	77.99	24.35
38	LEU	0	-34.92	-2E.97	79.01	26.24

39	LYS	N	-35.20	-25.71	77.17	26.25
39	LYS	CA	-36.59	-25.41	77.47	28.16
39	LYS	CB	-37.25	-24.61	76.34	28.65
39	LYS	CG	-38.35	-23.65	76.81	27.61
39	LYS	CD	-39.60	-24.37	77.25	27.20
39	LYS	CE	-40.68	-23.40	77.70	26.81
39	LYS	NZ	-41.94	-24.12	78.05	28.10
39	LYS	С	-37.37	-26.69	77.76	26.01
39	LYS	0	-38.28	-26.70	78.60	25.82
40	LYS	N	-37.00	-27.77	77.11	26.61
40	LYS	CA	-37.69	-29.03	77.34	27.03
40	LYS	CB	-37.64	-29.93	76.11	28.30
40	LYS	CG	-38.65	-31.06	76.16	29.91
40	LYS	CD	-38.79	-31.77	74.82	29.72
40	LYS	CE	-37.74	-32.83	74.61	26.31
40	LYS	NZ	-38.04	-33.61	73.37	31.45
40	LYS	С	-37.09	-29.72	78.56	25.36
40	LYS	0	-37.81	-30.32	79.35	23.91
41	LYS	N	-35.78	-29.57	78.73	25.39
41	LYS	CA	-35.06	-30.20	79.84	25.65
41	LYS	CB	-33.55	-30.06	79.66	24.57
41	LYS	CG	-32.72	-30.84	80.67	20.75
41	LYS	CD	-32.89	-32.34	80.50	26.88
41	LYS	CE	-31.76	-33.13	81.15	28.72
41	LYS	NZ	-31.63	-32.85	82.61	29.46
41	LYS	С	-35.50	-29.67	81.19	25.70
41	LYS	0	-35.92	-30.44	82.06	23.51
42	THR	N	-35.42	-28.35	81.34	26.91
42	THR	CA	-35.76	-27.67	82.58	25.94
42	THR	CB	-34.61	-26.77	83.03	26.05
42	THR	OG1	-34.60	-25.58	82.23	27.18
42	THR	CG2	-33.28	-27.49	82.85	28.99
42	THR	C	-37.00	-26.78	82.52	25.00
42	THR	0	-37.57	-26.43	83.55	28.90
43	GLY	N	-37.38	-26.35	81.32	24.41
43	GLY	CA	-38.54	-25.48	81.19	21.01
43	GLY	С	-38.09	-2402	81.13	19.83
43	GLY	0	-38.92	-23.10	81.08	14.91
44	LYS	N	-36.78	-23.82	81.15	22.11
44	LYS	CA	-36.15	-22.50	81.10	24.14
44	LYS	CB	-35.06	-22.38	82.17	26.33
44	LYS	CG	-35.61	-22.19	83.60	25.46

44	LYS	CD	-34.54	-22.37	84.69	26.23
44	LYS	CE	-33.36	-21.39	84.56	26.78
44	LYS	NZ	-32.26	-21.89	83.66	25.35
44	LYS	С	-35.63	-22.16	79.70	22.98
44	LYS	0	-34.86	-22.92	79.12	25.51
45	LEU	N	-36.06	-21.02	79.17	20.83
45	LEU	CA	-35.68	-20.55	77.84	14.37
45	LEU	CB	-36.93	-20.19	77.03	6.57
45	LEU	CG	-36.73	-19.98	75.54	7.56
45	LEŲ	CD1	-36.34	-21.28	74.88	7.35
45	LEU	CD2	-38.01	-19.44	74.92	7.68
45	LEU	С	-34.73	-19.36	77.92	12.93
45	LEU	0	-35.17	-18.22	78.02	12.18
46	LEU	N	-33.43	-19.62	77.91	12.78
46	LEU	CA	-32.45	-18.55	77.99	16.27
46	LEU	CB	-31.57	-18.68	79.23	17.06
46	LEU	CG	-32.20	-18.84	80.61	22.73
46	LEU	CD1	-33.29	-17.80	80.80	27.00
46	LEU	CD2	-32.76	-20.25	80.75	24.49
46	LEU	С	-31.59	-18.55	76.75	19.40
46	LEU	0	-31.38	-19.61	76.15	22.78
47	ASN	N	-31.05	-17.39	76.39	19.21
47	ASN	CA	-30.18	-17.27	75.21	16.88
47	ASN	CB	-30.12	-15.82	74.70	17.63
47	ASN	CG	-31.41	-15.38	74.02	19.44
47	ASN	OD1	-32.47	-15.31	74.64	24.63
47	ASN	ND2	-31.33	-15.04	72.74	18.95
47	ASN	С	-28.78	-17.79	75.50	14.05
47	ASN	0	-28.08	-17.23	76.36	11.81
48	LEU	N	-28.39	-18.87	74.83	12.40
48	LEU	CA	-27.05	-19.46	74.97	13.22
48	LEU	CB	-27.02	-20.94	74.58	14.23
48	LEU	CG	-27.64	-22.02	75.50	16.25
48	LEU	CD1	-26.96	-22.02	76.87	20.21
48	LEU	CD2	-29.13	-21.80	75.67	14.71
48	LEU	С	-26.08	-18.63	74.14	13.45
48	LEU	0	-26.51	-17.85	73.29	17.95
49	SER	N	-24.78	-18.80	74.36	15.33
49	SER	CA	-23.79	-17.98	73.65	13.70
49	SER	CB	-22.77	-17.41	74.65	14.72
49	SER	OG	-21.73	-16.68	74.02	13.12
49	SER	С	-23.05	-18.50	72.42	14.35

49	SER	0	-22.14	-19.33	72.52	15.35
50	PRO	N	-23.38	-17.93	71.24	13.10
50	PRO	CD	-24.49	-16.99	71.00	13.10
50	PRO	CA	-22.73	-18.30	69.99	8.82
50	PRO	CB	-23.41	-17.39	68.98	10.23
50	PRO	CG	-24.79	-17.23	69.54	10.20
50	PRO	С	-21.25	-17.95	70.09	7.83
50	PRO	0	-20.40	-18.65	69.56	7.76
51	GLN	N	-20.96	-16.86	70.80	8.14
51	GLN	CA	-19.59	-16.38	70.97	8.89
51	GLN	CB	-19.58	-14.99	71.62	12.14
51	GLN	CG	-18.31	-14.16	71.33	13.72
51	GLN	CD	-18.18	-13.71	69.87	12.95
51	GLN	OE1	-19.12	-13.18	69.27	10.44
51	GLN	NE2	-16.99	-13.89	69.30	6.01
51	GLN	С	-18.70	-17.34	71.74	8.59
51	GLN	0	-17.49	-17.46	71.45	8.17
52	asn	N	-19.27	-18.03	72.72	6.61
52	ASN	CA	-18.50	-19.01	73.49	6.19
52	ASN	CB	-19.36	-19.58	74.62	6.41
52	ASN	CG	-18.70	-20.74	75.34	7.63
52	ASN	OD1	-19.39	-21.64	75.81	9.07
52	ASN	ND2	-17.39	-20.70	75.48	4.94
52	ASN	C	-18.04	-20.09	72.50	9.66
52	ASN	0	-16.89	-20.56	72.54	9.03
53	LEU	N	-18.94	-20.43	71.58	10.50
53	LEU	CA	-18.69	-21.42	70.54	7.20
53	LEU	CB	-20.01	-21.83	69.89	6.76
53	LEU	CG	-20.74	-23.04	70.49	12.00
53	LEU	CD1	-20.47	-23.19	71.98	14.61
53	LEU	CD2	-22.22	-22.93	70.21	8.55
53	LEU	С	-17.69	-20.92	69.49	6.03
53	LEU		-16.75	-21.63	69.14	4.01
54	VAL	N	-17.87	-19.70	69.01	2.00
54	VAL	CA	-16.99	-19.14	68.00	6.81
54	VAL	CB	-17.34	-17.67	67.64	7.05
54	VAL	CG1	-16.27	-17.07	66.74	13.23
	VAL		-18.66	-17.58	66.93	5.36
54	VAL	C	-15.55	-19.15	68.47	12.15
54	VAL	0	-14.66	-19.63	67.76	16.44
55	ASP	N		-18.59	69.65	15.93
55	ASP	CA	-13.98	-18.48	70.22	17.26

55	ASP	CB	-13.98	-17.47	71.37	19.77
55	ASP	CG	-14.33	-16.06	70.94	22.96
55	ASP	OD1	-14.48	-15.78	69.72	24.84
55	ASP	OD2	-14.44	-15.21	71.84	23.39
55	ASP	С	-13.37	-19.78	70.72	18.42
55	ASP	0	-12.18	-20.06	70.49	12.75
56	CYS	N	-14.18	-20.55	71.43	20.14
56	CYS	CA	-13.75	-21.79	72.06	19.35
56	CYS	С	-13.66	-23.12	71.30	16.79
56	CYS	0	-13.00	-24.05	71.77	19.07
56	CYS	CB	-14.52	-21.96	73.37	17.64
56	CYS	SG	-14.48	-20.45	74.39	15.52
57	VAL	N	-14.31	-23.23	70.14	14.99
57	VAL	CA	-14.24	-24.47	69.38	12.09
57	VAL	СВ	-15.48	-24.72	68.51	9.25
57	VAL	CG1	-15.37	~26.09	67.85	2.60
57	VAL	CG2	-16.75	-24.62	69.34	5.09
57	VAL	С	-13.02	-24.48	68.47	12.69
57	VAL	0	÷13.04	-23.91	67.39	15.30
58	SER	N	-11.96	-25.14	68.92	15.12
58	SER	CA	-10.72	-25.21	68.15	18.94
58	SER	CB	-9.59	-25.78	69.00	21.08
58	SER	OG	-9.92	-27.07	69.49	20.89
58	SER	С	-10.84	-25.98	66.83	19.38
58	SER	0	-10.00	-25.81	65.94	21.21
59	GLU	N	-11.83	-26.87	66.74	21.74
59	GLU	CA	-12.04	-27.68	65.53	20.63
59	GLU	CB	-12.88	-28.94	65.82	20.59
59	GLU	CG	-12.48	-29.76	67.06	21.89
59	GLU	CD	-13.08	-29.21	68.37	21.83
59	GLU	OE1	-14.32	-29.11	68.47	25.51
59	GLU	OE2	-12.31	-28.90	69.30	26.68
59	GLU	С	-12.69	-26.87	64.41	19.86
59	GLU	0	-12.56	-27.21	63.23	21.27
60	ASN	N	-13.43	-25.83	64.79	16.86
60	ASN	CA	-14.11	-24.97	63.83	10.64
60	ASN	CB	-15.49	-24.61	64.33	2.00
60	ASN	CG	-16.43	-25.78	64.27	6.50
60	ASN	OD1	-17.46	-25.83	64.96	3.71
60	ASN	ND2	-16.09	-26.75	63.42	3.21
60	ASN			-23.76		13.09
	ASN		-12.25		64.13	14.35

61	ASP	N	-13.67	-23.01	62.46	14.85
61	ASP	CA	-12.90	-21.86	62.02	15.28
61	ASP	CB	-12.74	-21.86	60.51	21.27
61	ASP	CG	-11.30	-22.08	60.07	23.24
61	ASP	OD1	-10.60	-21.08	59.84	26.41
61	ASP	OD2	-10.88	-23.25	59.97	20.64
61	ASP	С	-13.36	-20.50	62.51	15.18
61	ASP	0	-12.89	-19.47	62.01	13.47
62	GLY	N	-14.26	-20.50	63.48	15.64
62	GLY	CA	-14.75	-19.25	64.03	15.74
62	GLY	С	-15.68	-18.55	63.07	17.26
62	GLY	0	-16.72	-19.10	62.72	19.72
63	CYS	N	-15.30	-17.35	62.64	19.00
63	CYS	CA	-16.14	-16.62	61.70	18.17
63	CYS	С	-16.00	-17.19	60.29	19.74
63	CYS	0	-16.79	-16.86	59.41	22.59
63	CYS	CB	-15.85	-15.11	61.73	16.37
63	CYS	SG	-16.32	-14.21	63.25	14.06
64	GLY	N	-15.00	-18.05	60.09	17.99
64	GLY	CA	-14.81	-18.67	58.79	15.84
64	GLY	C	-15.66	-19.92	58.60	17.59
64	GLY	0	-15.50	-20.66	57.63	19.05
65	GLY	N	-16.59	-20.15	59.53	18.26
65	GLY	CA	-17.46	-21.31	59.44	12.24
65	GLY	С	-17.03	-22.45	60.34	10.78
65	GLY	0	-15.90	-22.47	60.83	9.37
66	GLY	N	-17.94	-23.40	60.53	8.77
66	GLY	CA	-17.68	-24.56	61.36	8.46
66	GLY	С	-18.83	-25.55	61.24	10.14
66	GLY	0	-19.79	-25.29	60.51	9.60
67	TYR	N	-18.69	-26.70	61.88	12.33
67	TYR	CA	-19.74	-27.73	61.84	14.13
67	TYR	CB	-19.18	-29.12	61.49	14.00
67	TYR	CG	-18.53		60.13	16.82
67	TYR	CD1	-19.29	-29.29	58.96	19.29
67	TYR	CE1	-18.68	-29.37	57.70	19.34
67	TYR	CD2	-17.15	-29.24	60.02	17.85
67	TYR	CE2	16.53	-29.32	58.77	21.15
67	TYR	CZ	-17.30	-29.39	57.61	22.41
67	TYR	OH	-16.66	-29.42	56.38	22.94
67	TYR	С	-20.44	-27.77	63.20	12.95
67	TYR	0	-19.80	-27.61	64.23	14.93

68	MET N	-21.75	-27.98	63.19	14.10
68	MET CA	-22.51	-28.01	64.44	11.02
68	MET CB	-24.02	-28.04	64.18	7.28
68	MET CG	-24.57	-26.75	63.53	12.57
68	MET SD	-24.82	-26.81	61.72	9.43
68	MET CE	-23.29	-26.23	61.15	5.50
68	MET C	-22.07	-29.15	65.34	10.25
68	MET O	-21.98	-28.98	66.55	9.78
69	THR N	-21.74	-30.30	64.76	11.11
69	THR CA	-21.31	-31.45	65.54	9.38
69	THR CB	-21.06	-32.69	64.65	7.39
69	THR OG1	-20.19	-32.36	63.55	6.10
69	THR CG2	-22.39	-33.24	64.11	6.49
69	THR C	-20.08	-31.11	66.39	8.98
69	THR O	-20.06	-31.41	67.57	15.86
70	asn n	-19.09	-30.44	65.81	5.69
70	ASN CA	-17.89	-30.06	66.55	7.38
70	ASN CB	-16.87	-29.34	65.67	9.34
70	ASN CG	-16.43	-30.17	64.49	11.35
70	ASN OD1	-16.13	-29.63	63.43	13.75
70	ASN ND2	-16.38	-31.48	64.66	13.26
70	ASN C	-18.26	-29.15	67.71	9.60
70	ASN O	-17.68	-29.25	68.79	10.75
71	ALA N	-19.24	-28.28	67.47	12.80
71		-19.72	-27.33	68.48	15.69
71		-20.62	-26.29	67.84	12.87
71	ALA C	-20.48	-28.04	69.59	16.01
71	ALA O	-20.50	-27.56	70.72	18.32
72		-21.14	-29.14	69.25	15.89
72		-21.91	-29.90	70.23	15.72
72	PHE CB	-22.88	-30.85	69.53	12.85
72	PHE CG	-24.07	-30.17	68.91	10.89
72	PHE CD1	-24.61	-29.03	69.48	13.23
	PHE CD2	-24.69	-30.70	67.78	13.69
	PHE CE1	-25.75	-28.43	68.93	14.17
72		-25.83	-30.10	67.23	11.57
72		-26.36	-28.96	67.80	8.27
72	PHE C	-20.93	-30.65	71.11	17.93
72	PHE O	-20.98	-30.59	72.34	16.27
73	GLN N	-20.01	-31.36	70.44	19.40
73	GLN CA	-18.97	-32.16	71.08	19.37
73	GLN CB	-18.09	-32.78	70.00	17.22

.73	GLN	CG	-16.99	-33.70	70.46	19.79
73	GLN	CD	-16.30	-34.36	69.27	20.51
73	GLN	OE1	-16.73	-35.40	68.78	22.61
73	GLN	NE2	-15.24	-33.72	68.75	18.71
73	GLN	С	-18.15	-31.29	72.03	20.01
73	GLN	0	-17.74	-31.75	73.10	20.37
74	TYR	N	-17.92	-30.04	71.65	18.39
74	TYR	CA	-17.16	-29.12	72.48	15.45
74	TYR	СВ	-16.90	-27.79	71.73	13.54
74	TYR	CG	-16.82	-26.58	72.63	12.20
74	TYR	CD1	-15.63	-26.22	73.26	10.63
74	TYR	CE1	-15.59	-25.18	74.18	11.99
74	TYR	CD2	-17.97	-25.84	72.93	9.97
74	TYR	CE2	-17.94	-24.79	73.84	7.22
74	TYR	CZ	-16.75	-24.47	74.47	11.24
74	TYR	OH	-16.74	-23.48	75.43	17.85
74	TYR	C	-17.93	-28.93	73.80	12.38
74	TYR	0	-17.38	-29.14	74.87	15.74
75	VAL	N	-19.21	-28.60	73.71	11.39
75	VAL	CA	-20.05	-28.38	74.88	11.24
75	VAL	CB	-21.50	-28.06	74.44	8.02
75	VAL	CG1	-22.42	-27.95	75.64	6.92
75	VAL	CG2	-21.54	-26.76	73.66	9.76
75	VAL	C	-20.05	-29.59	75.83	16.34
75	VAL	0	-20.35	-29.45	77.03	11.43
76	GLN	N	-19.74	-30.76	75.28	18.49
76	GLN	CA	-19.69	-32.02	76.03	17.15
76	GLN	CB	-19.98	-33.22	75.11	16.55
76	GLN		-19.90	-34.60	75.78	17.80
76	GLN	CD	-20.05	-35.78	74.81	16.42
76	GLN	OE1	-19.33	-35.89	73.81	12.60
76	GLN		-20.98	-36.68	75.12	11.09
76	GLN		-18.33	-32.15	76.71	17.72
	GLN		-18.25	-32.28	77.92	19.51
	LYS		-17.25		75.94	18.51
	LYS		-15.91	-32.16	76.50	18.56
	LYS		-14.84	-32.18	75.41	18.21
77	LYS		-14.75	-33.47	74.62	18.16
	LYS		-13.48	-33.47	73.77	21.11
	LYS		-13.38	-34.69	72.84	22.51
	LYS		-12.02	-34.81	72.23	21.73
7 <b>7</b>	LYS	C	-15.58	-31.08	77.52	20.04

78 ASN N						
78 ASN CA	77	LYS O	-14.76	-31.29	78.42	21.61
78 ASN CB	78	B ASN N	-16.20	-29.91	77.36	20.45
78 ASN CG	78	asn ca	-15.96	-28.79	78.27	19.68
78 ASN OD1	78	ASN CB	-15.88	-27.47	77.50	18.78
78 ASN ND2	78	ASN CG	-15.19	-26.36	78.30	19.30
78 ASN C	78	ASN OD1	-13.97	-26.34	78.44	16.41
78 ASN C       -17.03       -28.73       79.35       18.44         78 ASN O       -17.03       -27.83       80.18       20.10         79 ARG N       -17.96       -29.68       79.32       16.26         79 ARG CA       -19.04       -29.74       80.30       16.53         79 ARG CB       -18.53       -30.31       81.63       16.34         79 ARG CG       -17.65       -31.54       81.47       17.65         79 ARG NE       -16.09       -31.93       82.78       19.26         79 ARG NE       -16.03       -33.01       82.58       21.25         79 ARG NE       -16.03       -33.01       82.58       21.25         79 ARG NH1       -14.27       -31.59       82.11       27.69         79 ARG NH2       -13.93       -33.86       82.13       18.73         79 ARG O       -19.74       -28.39       80.51       14.94         80 GLY N       -20.12       -27.73       79.42       16.45         80 GLY CA       -20.82       -26.46       79.56       14.39         81 ILE N       -21.64       -24.54       78.38       9.15         81 ILE CA       -21.62       -23.39       7	78	ASN ND2	-15.98	-25.40	78.79	20.55
78 ASN O	78	ASN C	-17.03	-28.73	79.35	18.44
79 ARG N -17.96 -29.68 79.32 16.26 79 ARG CA -19.04 -29.74 80.30 16.53 79 ARG CB -18.53 -30.31 81.63 16.34 79 ARG CG -17.65 -31.54 81.47 17.65 79 ARG CD -16.99 -31.93 82.78 19.26 79 ARG NE -16.03 -33.01 82.58 21.25 79 ARG NE -16.03 -33.01 82.58 21.25 79 ARG NE -16.03 -33.01 82.58 21.25 79 ARG NE -14.74 -32.82 82.27 24.36 79 ARG NH1 -14.27 -31.59 82.11 27.69 79 ARG NH2 -13.93 -33.86 82.13 18.73 79 ARG NH2 -13.93 -33.86 82.13 18.73 79 ARG O -19.91 -27.95 81.64 14.77 80 GLY N -20.12 -27.73 79.42 16.45 80 GLY C -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CA -21.62 -23.39 77.50 7.25 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 83 SER N -24.03 -18.28 78.87 11.91	78	ASN O	-17.03	-27.83	80.18	20.10
79 ARG CA -19.04 -29.74 80.30 16.53 79 ARG CB -18.53 -30.31 81.63 16.34 79 ARG CG -17.65 -31.54 81.47 17.65 79 ARG CD -16.99 -31.93 82.78 19.26 79 ARG NE -16.03 -33.01 82.58 21.25 79 ARG CZ -14.74 -32.82 82.27 24.36 79 ARG NH1 -14.27 -31.59 82.11 27.69 79 ARG NH2 -13.93 -33.86 82.13 18.73 79 ARG C -19.74 -28.39 80.51 14.94 79 ARG O -19.91 -27.95 81.64 14.77 80 GLY N -20.12 -27.73 79.42 16.45 80 GLY C -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 80 GLY O -19.49 -25.18 78.00 10.43 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CA -21.62 -23.39 77.50 7.25 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CB -19.95 -19.18 79.72 10.55 82 ASP CD -18.81 -18.86 79.31 8.90 82 ASP CD -23.46 -19.26 78.19 12.50 82 ASP C -23.46 -19.26 78.19 12.50 83 SER N -24.03 -18.28 78.87 11.91	79	ARG N	-17.96	-29.68	79.32	16.26
79 ARG CB	79	ARG CA	-19.04	-29.74	80.30	
79 ARG CG	79	ARG CB	-18.53	-30.31	81.63	
79 ARG CD	79	ARG CG	-17.65	-31.54		
79 ARG NE -16.03 -33.01 82.58 21.25 79 ARG CZ -14.74 -32.82 82.27 24.36 79 ARG NH1 -14.27 -31.59 82.11 27.69 79 ARG NH2 -13.93 -33.86 82.13 18.73 79 ARG C -19.74 -28.39 80.51 14.94 79 ARG O -19.91 -27.95 81.64 14.77 80 GLY N -20.12 -27.73 79.42 16.45 80 GLY CA -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 80 GLY O -19.49 -25.18 78.00 10.43 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CA -21.62 -23.39 77.50 7.25 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG -22.39 -23.68 76.48 4.07 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CG1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	79	ARG CD	-16.99	-31.93	82.78	19.26
79 ARG CZ -14.74 -32.82 82.27 24.36 79 ARG NH1 -14.27 -31.59 82.11 27.69 79 ARG NH2 -13.93 -33.86 82.13 18.73 79 ARG C -19.74 -28.39 80.51 14.94 79 ARG O -19.91 -27.95 81.64 14.77 80 GLY N -20.12 -27.73 79.42 16.45 80 GLY CA -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 80 GLY O -19.49 -25.18 78.00 10.43 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG2 -23.87 -23.89 76.48 4.07 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	79	ARG NE	-16.03	-33.01	82.58	21.25
79 ARG NH2   -13.93   -33.86    82.13    18.73 79 ARG C    -19.74   -28.39    80.51    14.94 79 ARG O    -19.91   -27.95    81.64    14.77 80 GLY N	79	ARG CZ	-14.74	-32.82	82.27	24.36
79 ARG NH2 -13.93 -33.86 82.13 18.73 79 ARG C -19.74 -28.39 80.51 14.94 79 ARG O -19.91 -27.95 81.64 14.77 80 GLY N -20.12 -27.73 79.42 16.45 80 GLY CA -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 80 GLY O -19.49 -25.18 78.00 10.43 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CA -21.62 -23.39 77.50 7.25 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG -22.39 -23.68 76.19 6.85 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE O -23.29 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	79	ARG NH1	-14.27	-31.59	82.11	27.69
79 ARG O -19.91 -27.95 81.64 14.77 80 GLY N -20.12 -27.73 79.42 16.45 80 GLY CA -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 80 GLY O -19.49 -25.18 78.00 10.43 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CA -21.62 -23.39 77.50 7.25 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG2 -23.87 -23.89 76.48 4.07 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CG1 -23.07 -22.63 73.97 11.83 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE O -23.29 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	79	ARG NH2	-13.93	-33.86	82.13	18.73
80 GLY N -20.12 -27.73 79.42 16.45 80 GLY CA -20.82 -26.46 79.56 14.39 80 GLY C -20.59 -25.34 78.56 13.10 80 GLY O -19.49 -25.18 78.00 10.43 81 ILE N -21.64 -24.54 78.38 9.15 81 ILE CA -21.62 -23.39 77.50 7.25 81 ILE CB -22.39 -23.68 76.19 6.85 81 ILE CG2 -23.87 -23.89 76.48 4.07 81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE O -23.29 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CB -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	79	ARG C	-19.74	-28.39	80.51	14.94
80 GLY CA	79	ARG O	-19.91	-27.95	81.64	14.77
80 GLY C	80	GLY N	-20.12	-27.73	79.42	16.45
80 GLY O	80	GLY CA	-20.82	-26.46	79.56	14.39
81 ILE N	80	GLY C	-20.59	-25.34	78.56	13.10
81 ILE CA	80	GLY O	-19.49	-25.18	78.00	10.43
81 ILE CB	81	ILE N	-21.64	-24.54	78.38	9.15
81 ILE CG2	81	ILE CA	-21.62	-23.39	77.50	7.25
81 ILE CG1 -22.16 -22.55 75.18 8.21 81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE O -23.29 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CB -19.95 -19.18 79.72 10.55 82 ASP CD -18.81 -18.86 79.31 8.90 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	81	ILE CB	-22.39	-23.68	76.19	6.85
81 ILE CD1 -23.07 -22.63 73.97 11.83 81 ILE C -22.27 -22.21 78.25 8.64 81 ILE O -23.29 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	81	ILE CG2	-23.87	-23.89	76.48	4.07
81 ILE C -22.27 -22.21 78.25 8.64 81 ILE O -23.29 -22.38 78.94 9.51 82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	81	ILE CG1	-22.16	-22.55	75.18	8.21
81 ILE O	81	ILE CD1	-23.07	-22.63	73.97	11.83
82 ASP N -21.68 -21.03 78.13 11.32 82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	81	ILE C	-22.27	-22.21	78.25	8.64
82 ASP CA -22.20 -19.84 78.81 12.61 82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	81	ILE O	-23.29	-22.38	78.94	9.51
82 ASP CB -21.14 -18.75 78.92 10.86 82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91			-21.68	-21.03	78.13	11.32
82 ASP CG -19.95 -19.18 79.72 10.55 82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	82	ASP CA	-22.20	-19.84	78.81	12.61
82 ASP OD1 -18.81 -18.86 79.31 8.90 82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	82	ASP CB	-21.14	-18.75	78.92	10.86
82 ASP OD2 -20.14 -19.86 80.74 14.89 82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	82	ASP CG	-19.95	-19.18	79. <b>7</b> 2	10.55
82 ASP C -23.46 -19.26 78.19 12.50 82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	82	ASP OD1	-18.81	-18.86	79.31	8.90
82 ASP O -23.92 -19.70 77.14 15.36 83 SER N -24.03 -18.28 78.87 11.91	82	ASP OD2	-20.14	-19.86	80.74	14.89
83 SER N -24.03 -18.28 78.87 11.91	82	ASP C	-23.46	-19.26	78.19	12.50
	82	ASP O	-23.92	-19.70	77.14	15.36
83 SER CA -25.23 -17.61 78.39 12.55	83	SER N	-24.03	-18.28	78.87	11.91
	83	SER CA	-25.23	-17.61	78.39	12.55

83	SER	CB	-26.14	-17.24	79.55	18.80
83	SER	OG	-25.48	-16.36	80.45	22.70
83	SER	С	-24.78	-16.35	77.64	9.20
83	SER	0	-23.59	-16.01	77.64	5.74
84	GLU	N	-25.71	-15.67	76.99	9.19
84	GLU	CA	-25.33	-14.43	76.31	14.72
84	GLU	CB	-26.47	-13.88	75.45	13.64
84	GLU	CG	-26.29	-14.13	73.96	13.53
84	GLU	CD	-25.09	-13.41	73.36	12.40
84	GLU	OE1	-24.88	-12.23	73.68	14.45
84	GLU	OE2	-24.38	-14.02	72.56	16.00
84	GLU	С	-24.86	-13.42	77.35	15.76
84	GLU	0	-23.85	-12.75	77.16	15.77
85	ASP	N	-25.57	-13.36	78.49	18.15
85	ASP	CA	-25.21	-12.45	79.58	18.14
85	ASP	CB	-26.05	-12.73	80.84	21.42
85	ASP	CG	-27.47	-12.20	80.72	27.07
85	ASP	OD1	-28.43	-12.97	80.98	25.69
85	ASP	OD2	-27.63	-11.01	80.38	28.79
85	ASP	С	-23.73	-12.56	79.90	20.03
85	ASP	0	-22.98	-11.59	79.78	24.76
86	ALA	N	-23.32	-13.77	80.28	17.91
86	ALA	CA	-21.93	-14.08	80.62	16.67
86	ALA	CB	-21.87	-15.44	81.30	15.93
86	ALA	C	-20.99	-14.07	79.41	19.50
86	ALA	0	-19.76	-14.03	79.57	22.48
87	TYR	N	-21.56	-14.13	78.21	16.64
87	TYR	CA	-20.72	-14.15	77.01	16.44
87	TYR	CB	-20.31	-15.59	76.69	12.21
87	TYR	CG	-18.90	-15.72	76.20	5.93
87	TYR		-18.37	-14.81	75.28	6.36
87	TYR		-17.08	-14.96	74.80	2.57
87			-18.10	-16.78	76.62	7.25
87	TYR	CE2	-16.81	-16.93	76.14	3.11
	TYR	CZ	-16.31		75.23	2.00
87	TYR	OH	-15.04	-16.18	74.72	8.83
87	TYR	С	-21.41	-13.49	75.82	16.77
87	TYR	0	-21.87	-14.18	74.91	22.23
88	PRO		-21.48	-12.15	75.80	15.42
88	PRO		-20.91	-11.18	76.74	12.94
88			-22.13	-11.44	74.70	13.08
88	PRO	CB	-21.96	-9.97	75.08	12.38

88	PRO	CG	-21.82	-10.00	76.55	9.35
88	PRO	С	-21.49	-11.72	73.35	14.00
88	PRO	0	-20.33	-12.14	73.27	15.05
89	TYR	N	-22.25	-11.52	72.28	13.57
89	TYR	CA	-21.77	-11.73	70.93	14.96
89	TYR	CB	-22.90	-12.16	69.99	16.79
89	TYR	CG	-22.46	-12.46	68.56	19.46
89	TYR	CD1	-21.64	-13.54	68.28	17.96
89	TYR	CE1	-21.26	-13.85	66.98	20.62
89	TYR	CD2	-22.90	-11.67	67.49	17.49
89	TYR	CE2	-22.53	-11.96	66.17	17.03
89	TYR	CZ	-21.70	-13.06	65.92	17.44
89	TYR	OH	-21.28	-13.36	64.65	16.20
89	TYR	С	-21.11	-10.47	70.39	14.19
89	TYR	0	-21.79	-9.45	70.23	14.43
90	VAL	N	-19.80	-10.50	70.18	11.92
90	VAL	CA	-19.15	-9.33	69.61	12.81
90	VAL	CB	-17.69	-9.13	70.09	11.22
90	VAL	CG1	-17.68	-8.76	71.57	19.09
90	VAL	CG2	-16.84	-10.35	69.83	10.92
90	VAL	Ċ	-19.21	-9.39	68.09	14.23
90	VAL	0	-19.09	-8.36	67.42	14.19
91	GLY	N	-19.44	-10.58	67.53	14.09
91	GLY	CA	-19.52	-10.72	66.09	15.26
91	GLY	C	-18.16	-10.68	65.44	18.32
91	GLY	0	-18.01	-10.38	64.24	16.36
92	GLN	N	-17.16	-11.06	66.23	20.31
92	GLN	CA	-15.78	-11.06	65.81	20.17
92	GLN	CB	-15.16	-9.69	66.09	23.65
92	GLN	CG	-13.83	-9.44	65.42	28.99
92	GLN	CD	-13.36	-8.00	65.62	34.47
92	GLN	OE1	-14.10	-7.15	66.15	34.75
	GLN		-12.14	-7.72	65.19	37.63
92	GLN	С	-15.09	-12.17	66.60	20.89
92	GLN	0	-15.53	-12.53	67.70	18.43
93	GLU	N	-14.04	-12.73	66.03	22.82
93	GLU	CA	-13.31	-13.81	66.69	24.92
93	GLU	CB	-12.63	-14.74	65.66	27.98
	GLU	CG	-12.23	-16.10	66.22	29.38
93	GLU	CD	-12.00	-17.17	65.15	34.25
93	GLU	OE1	-12.30	-16.93	63.95	35.80
93	GLU	OE2	-11.52	-13.26	65.52	34.14

93	GLU	С	-12:31	-13.29	67.71	24.19
93	GLU	0	-11.61	-12.31	67.45	24.56
94	GLU	N	-12.28	~13.94	68.87	23.61
94	GLU	CA	-11.38	-13.58	69.96	24.32
94	GLU	CB	-12.02	-12.55	70.90	22.62
94	GLU	CG	-12.23	-11.20	70.27	23.90
94	GLU	CD	-12.86	-10.23	71.21	24.44
94	GLU	OE1	-12.18	-9.26	71.60	29.56
94	GLU	OE2	-14.04	-10.43	71.57	23.54
94	GLU	С	-10.99	-14.84	70.73	23.80
94	GLU	0	-11.64	-15.89	70.59	21.98
95	SER	N	-9.95	-14.73	71.55	20.75
95	SER	CA	-9.47	-15.86	72.36	19.71
95	SER	CB	-8.26	-15.46	73.19	20.00
95	SER	OG	-8.57	-14.41	74.09	24.28
95	SER	C	-10.60	-16.38	73.25	20.63
95	SER	0	-11.48	-15.60	73.65	22.33
96	CYS	N	-10.56	-17.66	73.58	17.74
96	CYS	CA	-11.60	-18.26	74.42	18.66
96	CYS	С	-11.59	-17.72	75.85	19.92
96	CYS	0	-10.58	-17.79	76.56	21.50
96	CYS	CB	-11.51	-19.78	74.41	16.39
96	CYS	SG	-12.75	-20.61	75.44	19.58
97	MET	N	-12.72	-17.14	76.26	17.89
97	MET	CA	-12.88	-16.59	77.60	18.67
97	MET	CB	-12.86	-15.07	77.57	17.60
97	MET	CG	-12.76	-14.43	78.94	18.29
97	MET	SD	-11.15	-13.66	79.17	26.40
97	MET	CE	-9.99	-15.01	78.76	19.16
97	MET	C	-14.18	-17.09	78.20	21.66
97	MET	0	-15.07	-16.31	78.52	25.61
98	TYR	N	-14.30	-18.41	78.29	23.47
	TYR		-15.49	-19.06	78.83	23.75
98	TYR	CB	-15.58	-20.50	78.30	21.70
	TYR			-21.46	79.13	18.51
98	TYR	CD1		-21.54	78.90	18.20
98	TYR	CE1		-22.49	79.70	19.99
98	TYR	CD2	-15.80	-22.16	80.17	16.39
98	TYR	CE2	-16.53	-23.01	80.97	12.68
98	TYR	CZ		-23.17	80.74	16.90
98	TYR	OH		-23.99	81.57	22.57
98	TYR	С	-15.48	-19.01	80.37	26.07

98	TYR	. 0	-14.44	-19.23	80.99	27.32
99	ASN		-16.63	-18.67	80.96	28.21
99	ASN	CA	-16.76	-18.59	82.42	27.40
99	ASN	CB	-17.24	-17.20	82.85	30.57
99	ASN	CG	-17.89	-17.21	84.22	33.86
99	ASN	OD1	-19.06	-16.84	84.36	33.41
99	ASN	ND2	-17.16	-17.66	85.24	34.39
99	asn	С	-17.67	-19.66	83.03	25.48
99	ASN	0	-18.88	-19.64	82.82	25.31
100	PRO	N	-17.12	-20.53	83.87	22.21
100	PRO	CD	-15.70	-20.60	84.27	21.06
100	PRO	CA	-17.89	-21.60	84.50	19.72
100	PRO	CB	-16.89	-22.18	85.49	21.18
100	PRO	CG	-15.58	-22.01	84.75	22.13
100	PRO	C	-19.14	-21.08	85.22	19.80
100	PRO	0	-20.22	-21.65	85.0 <del>9</del>	15.82
101	THR	N	-18.98	-19.97	85.92	22.73
101	THR	CA	-20.07	-19.36	86.68	23.85
101	THR	CB	-19.53	-18.25	87.60	22.86
101	THR	OG1	-18.13	-18.47	87.87	20.58
101	THR	CG2	-20.28	-18.26	88.92	21.48
101	THR	C	-21.22	-18.80	85.82	24.75
101	THR	0	-22.30	-18.46	86.33	22.15
102	GLY	N	-20.96	-18.70	84.51	25.15
102	GLY	CA	-21.96	-18.20	83.59	21.81
102	GLY		-22.72	-19.34	82.94	19.11
102	GLY	0	-23.78	-19.12	82.36	19.57
103	LYS	N	-22.15	-20.55	83.03	18.20
103	LYS	CA	-22.73	-21.77	82.47	16.81
103	LYS	CB	-22.08	-23.02	83.07	16.94
103	LYS	CG	-22.90	-24.30	82.93	15.32
103	LYS		-22.03	-25.53	83.10	18.87
	LYS		-21.32	-25.57	84.46	21.19
103	LYS	NZ	-20.02	-26.34	84.40	20.74
	LYS	С	-24.24	-21.83	82.62	17.92
103	LYS		-24.78	-21.68	83.72	18.81
	ALA		-24.92	-22.05	81.50	17.95
	ALA		-26.38	-22.11	81.49	17.78
	ALA		-26.93	-21.14	80.47	21.54
104	ALA	С		-23.52	81.18	18.07
	ALA			-23.88	81.47	16.81
105	ALA	N	-25.95	-24.31	80.58	15.87

105	ALA	CA	-26.27	-25.69	80.23	13.04
105	ALA	CB	-27.41	-25.72	79.22	10.01
105	ALA	С	-25.05	-26.40	79.66	11.95
105	ALA	0	-23.94	-25.86	79.68	8.46
106	LYS	N	-25.27	-27.63	79.20	12.25
106	LYS	CA	-24.24	-28.45	78.57	18.76
106	LYS	CB	-23.26	-29.07	79.58	23.73
106	LYS	CG	-23.89	-30.02	80.60	26.39
106	LYS	CD	-22.88	-31.05	81.09	29.44
106	LYS	CE	-23.54	-32.05	82.06	33.28
106	LYS	NZ	-24.76	-32.74	81.51	24.10
106	LYS	С	-24.95	-29.54	77.77	18.52
106	LYS	0	-26.18	-29.49	77.64	16.87
107	CYS	N	-24.21	-30.50	77.22	19.97
107	CYS	CA	-24.81	-31.59	76.45	21.83
107	CYS	CB	-25.13	-31.16	75.00	23.15
107	CYS	SG	-23.72	-30.88	73.87	21.75
107	CYS	C	-23.98	-32.87	76.48	22.08
107	CYS	0	-22.75	-32.84	76.60	21.41
108	ARG	N	-24.66	-34.00	76.40	22.61
108	ARG	CA	-23.96	-35.28	76.44	25.98
108	ARG	CB	-24.57	-36.18	77.53	25.00
108	ARG	CG	-24.76	-35.44	78.85	19.33
108	ARG	CD	-25.24	-36.32	79.99	17.89
108	ARG	NE	-26.61	-36.80	79.81	10.20
108	ARG	CZ	-27.00	-38.02	80.15	9.00
108	ARG	NH1	-26.12	-38.88	80.68	6.62
108	ARG	NH2	-28.26	-38.39	79.99	10.32
108	ARG	С	-23.91	-35.98	75.08	23.86
108	ARG	0	-24.38	-37.10	74.94	25.08
109	GLY	N	-23.34	-35.29	74.10	23.61
	GLY		-23.21	-35.84	72.76	20.68
	GLY		-24.30	-35.35	71.82	18.95
109	GLY	0	-25.15	-34.56	72.22	19.25
	TYR		-24.29		70.58	18.18
	TYR		-25.27		69.57	17.03
	TYR		-24.72	-34.38	68.64	14.76
110	TYR	CG	-23.48	-34.81	67.90	12.25
110	TYR	CD1		-34.48	68.37	14.76
110	TYR	CE1		-34.95	67.75	14.89
110	TYR	CD2			66.78	12.73
110	TYR	CE2	-22.41	-36.08	66.14	15.02

110	TYR	CZ	-21.17	-35.75	66.64	17.33
110	TYR	OH	-20.03	-36.26	66.03	22.26
110	TYR	С	-25.70	-36.67	68.75	16.81
110	TYR	0	-25.01	-37.69	68.73	19.27
111	ARG	N	-26.83	-36.53	68.07	16.77
111	ARG	CA	-27.34	-37.57	67.19	18.25
111	ARG	CB	-28.73	-38.08	67.60	20.39
111	ARG	CG	-29.51	-38.81	66.48	21.92
111	ARG	CD	-28.94	-40.20	66.16	26.51
111	ARG	NE	-29.54	-40.85	64.98	25.17
111	ARG	CZ	-28.83	-41.49	64.03	26.40
111	ARG	NH1	-27.50	-41.55	64.12	21.95
111	ARG	NH2	-29.45	-42.10	63.02	18.73
111	ARG	С	-27.43	-36.92	65.81	20.54
111	ARG	0	-27.88	-35.77	65.69	23.20
112	GLU	N	-26.95	-37.62	64.79	17.24
112	GLU	CA	-27.01	-37.12	63.42	13.69
112	GLU	CB	-25.70	-37.34	62.69	12.60
112	GLU	CG	-24.59	-36.38	63.05	10.04
112	GLU	CD	-23.25	-36.92	62.62	11.35
112	GLU	OE1	-22.50	-36.23	61.89	10.65
112	GLU	OE2	-22.97	-38.06	63.01	14.44
112	GLU	С	-28.15	-37.85	62.72	15.47
112	GLU	0	-28.49	-38.97	63.09	17.83
113	ILE	N	-28.72	-37.25	61.70	17.43
113	ILE	CA	-29.82	-37.87	60.96	14.32
113	ILE	CB	-30.99	-36.87	60.90	13.01
113	ILE	CG2	-32.04	-37.32	59.87	11.54
113	ILE	CG1	-31.55	-36.70	62.33	10.24
	ILE		-32.35	-35.44	62.57	7.36
113	ILE		-29.38	-38.34	59.57	12.75
113	ILE		-28.59	-37.68	58.91	11.81
	PRO		-29.83	-39.53	59.13	15.44
114	PRO	CD	-30.82	-40.40	59.80	19.88
•	PRO			-40.08	57.81	15.21
	PRO	CB	-30.54	-41.13	57.59	15.21
114	PRO		-30.79	-41.66	58.96	16.85
114	PRO		-29.53	-39.00	56.73	18.58
114	PRO		-30.52	-38.27	56.64	21.69
	GLU			-38.91	55.91	17.93
115	GLU			-37.90	54.87	18.15
115	GLU	CB	-27.12	-37.95	54.11	21.41

115	GLU	CG	-27.04	-39.05	53.06	28.55
115	GLU	CD	-25.73	-39.04	52.26	34.52
115	GLU	OE1	-25.24	-37.94	51.90	34.27
115	GLU	OE2	-25.21	-40.14	51.99	34.99
115	GLU	С	-29.63	-37.99	53.90	20.57
115	GLU	0	-30.10	-39.08	53.56	20.47
116	GLY	N	-30.13	-36.82	53.51	21.25
116	GLY	CA	-31.24	-36.73	52.58	19.05
116	GLY	С	-32.59	-37.14	53.12	19.97
116	GLY	0	-33.62	-36.74	52.56	24.15
117	ASN	N	-32.62	-37.87	54.23	17.37
117	ASN	CA	-33.90	-38.33	54.77	15.93
117	ASN	CB	-33.73	-39.57	55.64	10.26
117	ASN	CG	-35.03	-40.34	55.79	12.00
117	ASN	OD1	-36.12	~39.76	55.79	10.30
117	ASN	ND2	-34.92	-41.66	55.91	13.10
117	ASN	С	-34.73	-37.30	55.49	17.41
117	ASN	0	-34.57	-37.09	56.70	18.71
118	GLU	N	-35.66	-36.70	54.77	15.53
118	GLU	CA	-36.56	-35.71	55.34	15.33
118	GLU	CB	-37.27	-34.93	54.25	15.94
118	GLU	CG	-36.37	-34.00	53.46	22.20
118	GLU	CD	-37.14	-33.16	52.46	21.94
118	GLU	OE1	-37.27	-33.58	51.29	25.99
118	GLU	OE2	-37.64	-32.08	52.85	25.26
118	GLU	С	-37.57	-36.30	56.33	18.60
118	GLU	0	-38.02	-35.60	57.23	23.43
119	LYS	N	-37.93	-37.57	56.17	19.21
119	LYS	CA	-38.89	-38.21	57.08	17.16
119	LYS	CB	-39.39	-39.55	56.54	18.23
119	LYS	CG	-40.29	-39.44	55.34	23.03
119			-41.48	-38.55	55.62	24.88
119	LYS	CE	-41.85	-37.77	54.36	33.17
	LYS		-40.65	-37.07		35.51
119	LYS	C	-38.30		58.44	13.83
119	LYS	0	-38.99		59.45	14.48
120	ALA	N	-37.02		58.47	11.43
120			-36.30	-38.93	59.71	11.35
	ALA		-34.91	-39.44	59.43	12.94
	ALA				60.34	14.11
120	ALA	0	-36.49		61.53	20.36
121	LEU	N	-35.94	-36.57	59.52	14.09

121 LEU C	-35.84	-35.19	59.98	14.37
121 LEU CE	3 -35.35	-34.28	58.86	11.45
121 LEU CO	-35.11	-32.82	59.20	13.53
121 LEU CI	01 -34.22	-32.69	60.44	17.91
121 LEU CI	2 -34.49	-32.13	57.98	13.43
121 LEU C	-37.17	-34.71	60.57	14.14
121 LEU O	-37.19	-34.08	61.63	17.50
122 LYS N	-38.28	-35.01	59.90	14.51
122 LYS CA	-39.61	-34.63	60.36	14.97
122 LYS CE	-40.67	-34.97	59.32	15.25
122 LYS CO	-42.09	-34.67	59.75	15.73
122 LYS CD	-43.09	-35.54	59.00	15.63
122 LYS CE	-44.48	-35.36	59.56	22.12
122 LYS NZ	-45.45	-36.35	59.02	25.27
122 LYS C	-39.93	-35.32	61.69	17.92
122 LYS O	-40.58	-34.74	62.58	15.41
123 ARG N	-39.51	-36.57	61.80	20.19
123 ARG CA	-39.71	-37.36	63.00	24.53
123 ARG CB	-39.36	-38.82	62.76	26.04
123 ARG CG	-40.37	-39.57	61.92	29.94
123 ARG CD	-39.79	-40.87	61.39	33.27
123 ARG NE	-38.98	-41.56	62.39	38.93
123 ARG CZ	-37.67	-41.79	62.26	38.64
123 ARG NH	1 -37.03	-41.37	61.18	39.92
123 ARG NH	2 -37.00	-42.45	63.20	34.71
123 ARG C	-38.89	-36.77	64.14	25.64
123 ARG O	-39.33	-36.76	65.30	25.72
124 ALA N	-37.70	-36.26	63.80	23.74
124 ALA CA	-36.80	-35.64	64.77	25.05
124 ALA CB	-35.49	-35.23	64.09	22.46
124 ALA C	-37.48	-34.41	65.37	24.90
124 ALA O	-37.50	-34.23	66.59	25.58
125 VAL N	-38.02	-33.57	64.50	25.44
125 VAL CA	-38.69	-32.35	64.92	24.60
125 VAL CB	-39.08	-31.46	63.72	20.30
125 VAL CG	1 -39.85	-30.23	64.19	13.27
125 VAL CG	2 -37.80	-31.05	62.95	19.58
125 VAL C	-39.93	-32.66	65.73	24.17
125 VAL O	-40.19	-32.01	66.73	26.08
126 ALA N	-40.69	-33.66	65.32	25.26
126 ALA CA	-41.90	-34.00	66.05	26.85
126 ALA CB	-42.65	-35.08	65.34	27.49

126	ALA	С	-41.59	-34.41	67.50	28.51
126	ALA	0	-42.22	-33.91	68.43	29.45
127	ARG	N	-40.60	-35.30	67.68	27.15
127	ARG	CA	-40.20	-35.80	69.01	26.50
127	ARG	CB	-39.52	-37.17	68.92	25.80
127	ARG	CG	-40.36	-38.31	68.39	29.17
127	ARG	CD	-41.39	-38.82	69.38	30.20
127	ARG	NE	-42.01	-40.06	68.90	31.67
127	ARG	CZ	-43.14	-40.57	69.38	31.87
127	ARG	NH1	-43.62	-41.70	68.87	32.34
127	ARG	NH2	-43.80	-39.97	70.36	33.34
127	ARG	С	-39.30	-34.88	69.83	26.63
127	ARG	0	-39.61	-34.54	70.98	29.49
128	VAL	N	-38.13	-34.58	69.27	24.07
128	VAL	CA	-37.14	-33.75	69.93	20.27
128	VAL	CB	-35.74	-33.96	69.29	17.18
128	VAL	CG1	-34.73	-33.00	69.87	16.62
128	VAL	CG2	-35.27	-35.40	69.51	16.17
128	VAL	С	-37.51	-32.28	69.92	19.91
128	VAL	0	-38.00	-31.75	70.91	21.31
129	GLY	N	-37.26	-31.63	68.79	21.40
129	GLY	CA	-37.56	-30.22	68.67	18.08
129	GLY	С	-36.62	-29.60	67.67	17.76
129	GLY	0	-36.22 <sup>-</sup>	-30.27	66.71	14.46
130	PRO	N	-36.25	-28.33	67.86	18.11
130	PRO	CD	-36.68	-27.44	68.96	20.59
130	PRO	CA	-35.35	-27.64	66.94	17.36
130	PRO	CB	-35.03	-26.36	67.70	17.67
130	PRO	CG	-36.33	-26.07	68.41	19.12
130	PRO	С	-34.10	-28.47	66.69	18.18
130	PRO	0	-33.43	-28.92	67.63	21.49
131	VAL		-33.81	-28.71	65.41	14.71
	VAL		-32.66	-29.47	64.99	10.15
131	VAL					13.09
131	VAL	CG1	-31.88		63.55	14.17
131	VAL	CG2	-33.88		65.02	19.95
131	VAL	С		-28.60	64.10	10.57
	VAL	0	-32.33	-27.86	63.27	13.19
132	SER	N		-28.71	64.25	7.45
132	SER	CA		-27.93	63.42	7.72
132	SER	CB			64.07	9.68
132	SER	OG	-28.37	-27.45	65.44	20.96

. 132	SER	C	-29.48	-28.55	62.02	6.88
132	SER	. 0	-29.18	-29.74	61.86	4.58
133	VAL	N	-29.75	-27.75	61.00	7.92
133	VAL	CA	-29.68	-28.20	59.62	6.47
133	VAL	CB	-31.09	-28.35	59.00	6.27
133	VAL	CG1	-31.92	-29.35	59.80	5.31
133	VAL	CG2	-31.80	-27.00	58.93	2.32
133	VAL	С	-28.88	-27.20	58.78	6.11
133	VAL	0	-28.93	-25.99	59.01	4.92
134	ALA	N	-28.09	-27.71	57.85	8.55
134	ALA	CA	-27.29	-26.90	56.94	10.56
134	ALA	CB	-25.90	-27.49	56.80	10.48
134	ALA	С	-28.01	-26.89	55.59	11.28
134	ALA	0	-28.61	-27.91	55.19	10.70
135	ILE	N	-28.00	-25.75	54.91	5.87
135	ILE	CA	-28.65	-25.64	53.61	9.02
135	ILE	CB	-30.05	-25.04	53.75	9.82
135	ILE	CG2	-30.96	-25.98	54.52	8.18
135	ILE	CG1	-29.97	-23.67	54.42	7.95
135	ILE	CD1	-31.24	-22.87	54.30	5.36
135	ILE	С	-27.84	-24.75	52.67	10.58
135	ILE	0	-26.82	-24.17	53.08	11.83
	ASP		-28.27	-24.67	51.41	14.26
	ASP		-27.64	-23.83	50.39	12.77
	ASP		-27.59	-24.54	49.03	11.56
136	ASP	CG	-27.29	-23.58	47.87	11.70
136	ASP	OD1	-27.82	-23.80	46.77	16.82
	ASP		-26.54	-22.61	48.06	6.89
	ASP		-28.44	-22.53	50.29	11.76
	ASP		-29.51	-22.50	49.69	12.86
	ALA		-27.90	-21.45	50.83	12.10
	ALA		-28.63	-20.19	50.80	14.12
	ALA		-28.77	-19.64	52.22	16.96
	ALA		-28.02		49.87	15.39
	ALA		-28.27		50.04	11.83
	SER			-19.57	48.91	16.46
	SER		-26.56	-18.66	47.96	16.14
138	SER		-25.45	-19.36	47.17	13.20
138	SER			-20.60	46.65	11.80
138				-17.97	47.02	
138				-16.77		
139	LEU	N	-28.5C	-18.75	46.53	11.84

139	LEU	CA	-29.51	-18.24	45.62	10.88
139	LEU	CB	-30.62	-19.27	45.41	10.28
139	LEU	CG	-30.39	-20.25	44.25	7.79
139	LEU	CD1	-28.98	-20.81	44.28	16.09
139	LEU	CD2	-31.39	-21.35	44.31	6.66
139	LEU	С	-30.07	-16.90	46.07	10.95
139	LEU	0	-30.56	-16.77	47.18	10.84
140	THR	N	-29.98	-15.89	45.20	12.65
140	THR	CA	-30.48	-14.56	45.55	10.04
140	THR	CB	-30.04	-13.47	44.54	8.74
140	THR	OG1	-30.71	-13.68	43.29	16.41
140	THR	CG2	-28.52	-13.50	44.33	5.90
140	THR	С	-31.98	-14.51	45.76	7.53
140	THR	0	-32.51	-13.48	46.16	7.86
141	SER	N	-32.69	-15.60	45.48	7.62
141	SER	CA	-34.13	-15.60	45.71	10.59
141	SER	CB	-34.81	-16.69	44.90	7.78
141	SER	OG	-34.14	-17.92	45.09	10.19
141	SER	С	-34.32	-15.81	47.21	14.46
141	SER	0	-35.31	-15.38	47.80	15.07
142	PHE	N	-33.33	-16.47	47.81	16.29
142	PHE	CA	-33.32	-16.75	49.24	18.29
142	PHE	CB	-32.26	-17.80	49.57	19.91
142	PHE	CG	-32.23	-18.22	51.01	19.34
142	PHE	CD1	-32.83	-19.42	51.40	19.10
142	PHE	CD2	-31.58	-17.44	51.96	17.86
142	PHE	CE1	-32.77	-19.84	52.72	18.29
142	PHE		-31.52	-17.86	53.29	16.09
142	PHE	CZ	-32.12	-19.06	53.67	17.03
142	PHE	С	-33.04	-15.46	49.98	18.87
142	PHE	0	-33.70	-15.13	50.96	20.82
143			-32.06	-14.71	49.48	17.89
	GLN		-31.69	-13.46	50.11	17.94
143	GLN	CB			49.56	20.86
	GLN			-14.07	49.66	27.97
	GLN		-27.96	-13.66	49.06	35.25
	GLN		-27.88	-12.65	48.36	42.66
	GLN			-14.44	49.35	32.14
	GLN			-12.40	49.97	16.38
	GLN			-11.43	50.72	21.06
144	PHE	N	-33.71	-12.62	49.07	16.70
144	PHE	CA	-34.81	-11.68	48.86	16.93

144 PHE CB	-34.83	-11.16	47.42	14.35
144 PHE CG	-33.63	-10.34		9.94
144 PHE CD1	-33.23	-9.28		
144 PHE CD2	-32.87	-10.63	45.94	
144 PHE CE1	-32.09	-8.54	47.58	2.23
144 PHE CE2	-31.73	-9.89	45.63	11.16
144 PHE CZ	-31.34	-8.85	46.45	5.01
144 PHE C	-36.15	-12.29		19.07
144 PHE O	-37.21			23.82
145 TYR N	-36.10	-13.42	49.95	21.54
145 TYR CA	-37.31	-14.10	50.39	20.66
145 TYR CB	-36.98	-15.43	51.09	
145 TYR CG	-38.13	-16.04		20.41
145 TYR CD1	-38.96			22.31
145 TYR CE1	-40.05	-17.51	51.93	23.37
145 TYR CD2	-38.43	-15.64	53.15	19.98
145 TYR CE2	-39.52		53.84	20.51
145 TYR CZ	-40.33	-17.09	53.22	22.20
145 TYR OH	-41.42	-17.60	53.89	24.45
145 TYR C	-38.13	-13.18	51.31	19.31
145 TYR O	-37.57	-12.41	52.09	16.29
146 SER N	-39.44	-13.29	51.23	19.16
146 SER CA	-40.31	-12.46	52.05	18.38
146 SER CB	-40.69	-11.17	51.33	21.74
146 SER OG	-41.24	-11.45	50.05	25.62
146 SER C	-41.55	-13.21	52.52	17.33
146 SER O	-41.87	-13.19	53.70	17.00
147 LYS N	-42.21	-13.90	51.60	15.30
147 LYS CA	-43.42	-14.63	51.94	16.47
147 LYS CB	-44.65	-13.72	51.85	21.13
147 LYS CG	-45.98	-14.43	52.13	26.90
147 LYS CD	-47.15	-13.75	51.41	31.88
147 LYS CE	-48.38	-14.69	51.35	38.13
147 LYS NZ	-49.45	-14.23	50.41	36.35
147 LYS C	-43.61	-15.88	51.09	16.19
147 LYS O	-43.21	-15.91	49.92	18.85
148 GLY N	-44.20	-16.91	51.69	14.07
148 GLY CA	-44.47	-18.14	50.98	13.24
148 GLY C	-43.46	19.26	51.17	12.23
148 GLY O	-42.39	-19.05	51.74	11.43
149 VAL N	-43.82	-20.45	50.71	9.32
149 VAL CA	-42.95	-21.61	50.81	8.20

149 VAL	CB	-43.70	-22.96	50.60	5.83
149 VAL	CG1	-42.72	-24.11	50.73	2.00
149 VAL	CG2	-44.82	-23.12	51.61	4.34
149 VAL	C	-41.92	-21.43	49.70	11.54
149 VAL	0	-42.25	-21.45	48.50	12.48
150 TYR	. <b>N</b>	-40.68	-21.21	50.13	12.91
150 TYR	CA	-39.55	-21.02	49.24	10.42
150 TYR	CB	-38.33	-20.58	50.03	4.78
150 TYR	CG	-37.10	-20.42	49.19	8.23
150 TYR	CD1	-36.83	-19.21	48.55	10.09
150 TYR	CE1	-35.68	-19.06	47.77	11.54
150 TYR	CD2	-36.21	-21.47	49.02	8.80
150 TYR	CE2	-35.07	-21.33	48.24	9.65
150 TYR	CZ	-34.81	-20.13	47.62	10.30
150 TYR	OH	-33.67	-19.97	46.88	18.73
150 TYR	С	-39.27	-22.30	48.46	14.00
150 TYR	0	-39.45	-23.40	48.96	16.63
151 TYR	N	-38.79	-22.13	47.24	15.66
151 TYR		-38.48	-23.25	46.37	17.45
151 TYR	CB	-39.74	-24.04	46.01	21.54
151 TYR	CG	-39.53	-25.25	45.13	23.93
	CD1	-38.93	-26.42	45.64	28.17
	CE1	-38.74	-27.54	44.83	29.69
151 TYR	CD2	-39.94	-25.25	43.80	26.78
151 TYR	CE2	-39.76	-26.36	42.99	29.59
151 TYR		-39.16	-27.50	43.51	32.44
151 TYR	OH	-38.99	-28.60	42.70	37.61
151 TYR	С	-37.85	-22.60	45.15	18.03
151 TYR		-38.32	-21.56	44.68	16.75
152 ASP		-36.76	-23.17	44.68	19.75
152 ASP		-36.05	-22.64	43.52	21.69
152 ASP		-35.07	-21.53	43.91	21.35
152 ASP		-34.48	-20.80	42.70	22.73
152 ASP				42.63	20.20
152 ASP		-33.90		41.82	23.61
152 ASP		-35.33	-23.85	42.96	25.58
152 ASP		-34.67	-24.57	43.72	28.85
153 GLU			-24.09	41.66	26.65
153 GLU			<b>-25</b> .25	41.03	26.63
L53 GLU			-25.57	39.69	30.19
153 GLU			-24.38	38.92	34.99
L53 GLU	CD	-35.04	-23.61	38.15	37.99

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15			-22.38	38.34	40.14
15	3 GLU OE2	-34.31	-24.24	37.34	40.23
153	GLU C	-33.34	-25.20	40.91	24.39
153	GLU O	-32.71	-26.22	40.70	23.54
154	4 SER N	-32.75	-24.02	41.11	23.15
154	4 SER CA	-31.30	-23.83	41.04	25.05
154	SER CB	-30.95	-22.35	40.85	24.11
154	SER OG	-31.81	-21.71	39.91	26.07
154	SER C	-30.53	-24.38	42.26	25.62
154	SER O	-29.37	-24.78	42.15	26.42
155	CYS N	-31.19	-24.33	43.43	24.27
155	CYS CA	-30.63	-24.77	44.71	19.73
155	CYS C	-29.88	-26.08	44.59	17.05
155	CYS O	-30.44	-27.08	44.16	14.86
155	CYS CB	-31.72	-24.87	45.79	20.11
155	CYS SG	-31.27	-24.12	47.40	19.94
156	asn n	-28.61	-26.07	44.99	19.11
156	ASN CA	-27.76	-27.26	44.92	21.62
156	ASN CB	-26.37	-26.87	44.41	19.84
156	ASN CG	-25.47	-28.06	44.20	20.15
156	ASN OD1	-25.86	-29.20	44.44	15.60
156	ASN ND2	-24.23	-27.80	43.77	19.95
156	ASN C	-27.67	-28.05	46.24	22.71
156	ASN O	-27.08	-27.58	47.21	23.56
157	SER N	-28.20	-29.28	46.23	22.09
157	SER CA	-28.22	-30.17	47.40	16.83
157	SER CB	-28.93	-31.49	47.06	14.81
157	SER OG	-30.26	-31.26	46.66	19.44
157	SER C	-26.86	-30.49	48.02	17.43
157	SER O	-26.78	-31.10	49.10	16.02
158	ASP N	-25.79	-30.19	47.30	17.73
158	ASP CA	-24.46	-30.46	47.80	20.40
158	ASP CB	-23.70	-31.42	46.88	24.69
158	ASP CG	-24.27	-32.82	46.89	27.15
158	ASP OD1	-24.93	-33.19	45.88	26.00
158	ASP OD2	-24.08	-33.55	47.90	26.52
158	ASP C	-23.68	-29.19	48.00	20.27
158	ASP O	-22.46	-29.22	48.16	20.24
159	ASN N	-24.39	-28.07	47.97	17.11
159	ASN CA	-23.72	-26.81	48.19	16.88
159	ASN CB	-24.01	-25.79	47.10	18.24
159	ASN CG	-23.00	-24.56	47.10	21.17

159	ASN	OD1	-21.83	-24.86	47.42	16.98
159	ASN	ND2	-23.46	-23.46	46.75	22.38
159	ASN	С	-24.12	-26.28	49.55	17.56
159	ASN	0	-24.64	-25.17	49.67	19.13
160	LEU	N	-23.93	-27.12	50.57	12.99
160	LEU	CA	-24.23	-26.73	51.94	8.24
160	LEU	СВ	-24.00	-27.89	52.89	4.68
160	LEU	CG	-25.09	-28.95	53.11	5.66
160	LEU	CD1	-25.96	-29.16	51.89	3.96
160	LEU	CD2	-24.43	-30.24	53.54	6.24
160	LEU	С	-23.28	-25.58	52.23	8.29
160	LEU	0	-22.07	-25.75	52.11	8.75
161	ASN	N	-23.83	-24.41	52.54	9.39
161	ASN	CA	-23.02	-23.23	52.83	10.51
161	ASN	CB	-22.74	-22.43	51.56	14.18
	asn		-24.01	-22.01	50.85	10.89
161	asn	OD1	-24.80	-21.22	51.35	14.49
161	asn	ND2	-24.21	-22.56	49.67	16.31
161	ASN	С	-23.56	-22.29	53.92	12.43
	ASN	0	-23.02	-21.20	54.14	13.86
	HIS	N	-24.62	-22.69	54.60	11.27
	HIS		-25.19	-21.85	55.64	10.37
162	HIS	CB	-26.13	-20.82	55.00	5.69
162	HIS	CG	-26.80	-19.89	55.96	6.06
162	HIS	CD2	-28.10	<b>-19</b> .79	56.33	6.15
162	HIS	ND1	-26.14	-18.85	56.61	11.02
162	HIS	CE1	-27.01	-18.16	57.33	5.27
162		NE2	-28.20	-18.71	57.17	6.68
162	HIS		-25.92	-22.76	56.61	11.99
162	HIS		-26.68	-23.63	56.20	15.71
	ALA		-25.59	-22.65	57.89	14.09
	ALA		-26.22	-23.48	58.91	10.89
	ALA		-25.26	-23.79	60.01	12.33
	ALA			-22.67		8.08
	ALA			-21.48	59.69	7.93
	VAL			-23.33	59.61	8.61
	VAL			-22.67	60.06	9.19
	VAL			-22.36	58.82	13.42
	VAL			-23.51	58.50	8.48
	VAL			-21.00	58.93	12.71
	VAL			-23.60		9.94
164	VAL	0	-29.95	-24.69	61.35	11.85

165	LEU	N	-31.59	-23.20	61.61	5.74
165	LEU	CA	-32.29	-24.07	62.58	6.61
165	LEU	CB	-32.27	-23.43	63.97	3.31
165	LEU	CG	-32.85	-24.20	65.15	2.00
165	LEU	CD1	-31.81	-25.14	65.69	3.91
165	LEU	CD2	-33.27	-23.20	66.22	3.66
165	LEU	С	-33.71	-24.47	62.20	8.12
165	LEU	0	-34.52	-23.62	61.80	12.38
166	ALA	N	-34.04	-25.75	62.36	6.76
166	ALA	CA	-35.37	-26.25	62.02	6.92
166	ALA	CB	-35.28	-27.67	61.50	5.38
166	ALA	С	-36.29	-26.20	63.24	9.15
166	ALA	0	-36.23	-27.07	64.10	12.12
167	VAL	N	-37.15	-25.19	63.30	10.50
167	VAL	CA	-38.06	-25.06	64.42	10.11
167	VAL	CB	-38.16	-23.60	64.93	9.38
167	VAL	CG1	-36.79	-23.07	65.35	2.39
167	VAL	CG2	-38.82	-22.71	63.88	7.39
167	VAL	C	-39.47	-25.56	64.11	10.07
167	VAL	0	-40.44	-25.13	64.73	12.94
168	GLY	N	-39.60	-26.47	63.16	9.76
168	GLY	CA	-40.93	-26.97	62.85	7.51
168	GLY	С	-41.09	-27.55	61.46	8.86
168	GLY	0	-40.11	-27.88	60.80	7.84
169	TYR	N	-42.34	-27.71	61.06	10.03
169	TYR	CA	-42.70	-28.24	59.76	12.55
169	TYR	CB	-42.20	-29.67	59.56	12.27
169	TYR	CG	-42.79	-30.71	60.51	12.03
169	TYR	CD1	-44.13	-31.06	60.43	9.03
169	TYR	CE1	-44.67	-32.02	61.26	9.87
169	TYR	CD2	-42.00	-31.35	61.45	11.40
169	TYR	CE2	-42.54		62.30	8.77
169	TYR	CZ	-43.87	-32.64	62.19	6.67
169	TYR	OH	-44.45	-33.58	63.00	8.74
169	TYR	С	-44.21	-28.15	59.68	15.32
169	TYR	0	-44.88	-28.12	60.72	19.66
170	GLY	N	-44.77	-28.10	58.47	15.75
170	GLY	CA	-46.21	-28.00	58.36	13.53
170	GLY	С	-46.70	-28.03	56.94	14.99
1.70	GLY	0	-46.13	-28.72	56.09	14.31
171	ILE	N	-47.70	-27.21	56.66	17.47
171	ILE	CA	-48.28	-27.14	55.33	23.06

171	ILE	CB	-49.37	-28.23	55.17	25.40
171	ILE	CG2	-50.56	-27.93	56.06	30.02
171	ILE	CG1	-49.86	-28.31	53.73	33.16
171	ILE	CD1	-49.86	-29.73	53.19	37.53
171	ILE	С	-48.86	-25.75	55.07	21.85
171	ILE	0	-49.14	-25.02	56.01	26.83
172	GLN	N	-48.95	-25.33	53.81	24.21
172	GLN	CA	-49.52	-24.03	53.46	24.63
172	GLN	CB	-48.53	-22.88	53.66	22.98
172	GLN	CG	-49.13	-21.50	53.34	24.04
172	GLN	CD	-48.15	-20.35	53.49	25.53
172	GLN	OE1	-47.37	-20.30	54.45	30.09
172	GLN	NE2	-48.17	-19.42	52.55	28.42
172	GLN	С	-50.02	-24.03	52.02	25.83
172	GLN	0	-49.24	-24.28	51.09	28.71
173	LYS	N	-51.32	-23.80	51.83	23.80
173	LYS	CA	-51.91	-23.78	50.49	22.33
173	LYS	CB	-51.58	-22.46	49.78	22.26
	LYS	CG	-52.35	-21.23	50.29	26.56
173	LYS	CD	-53.81	-21.23	49.80	30.52
173	LYS	CE	-54.48	-19.85	49.89	29.44
173	LYS	NZ	-54.64	-19.30	51.27	28.52
173	LYS	С	-51.45	-24.98	49.66	21.89
173	LYS		-51.02	-24.84	48.52	22.66
174			-51.49	-26.16	50.27	21.87
174			-51.08	-27.38	49.61	20.01
174	GLY		-49.59	-27.70	49.71	19.12
174	GLY		-49.18	-28.85	49.49	14.08
	ASN		-48.79	-26.71	50.06	19.25
175	ASN		-47.34	-26.90	50.16	22.03
	ASN		-46.59	-25.65	49.70	23.43
175	ASN		-47.13		48.40	26.87
	ASN		-46.80	-25.56	47.31	24.31
	ASN			-24.04	48.52	27.46
	ASN			-27.30	51.56	22.36
	ASN				52.51	22.73
176				-28.52	51.69	18.82
	LYS		-45.79	-28.98	52.95	18.15
	LYS	CB	-45.64	-30.50	52.97	19.27
	LYS			-31.32	53.07	21.41
	LYS			-32.84		17.01
176	LYS	CE	-47.83	-33.72	53.27	21.32

176	LYS	NZ	-48.07	-33.97	54.73	17.44
176	LYS	С	-44.42	-28.30	53.03	20.70
176	LYS	0	-43.56	-28.53	52.17	21.72
177	HIS	N	-44.19	-27.51	54.08	18.91
177	HIS	CA	-42.93	-26.78	54.26	13.73
177	HIS	CB	-43.18	-25.28	54.19	16.90
177	HIS	CG	-44.04	-24.76	55.30	17.13
177	HIS	CD2	-43.77	-24.55	56.62	20.36
177	HIS	ND1	-45.37	-24.39	55.13	20.15
177	HIS	CE1	-45.87	-23.99	56.28	21.57
177	HIS	NE2	-44.92	-24.08	57.20	21.14
177	HIS	С	-42.20	-27.08	55.56	13.57
177	HIS	0	-42.71	-27.80	56.42	15.98
178	TRP	N	-41.05	-26.45	55.74	11.27
178	TRP	CA	-40.27	-26.56	56.97	10.08
178	TRP	CB	-38.80	-26.95	56.72	13.29
178	TRP	CG	-38.53	-28.36	56.29	9.82
178	TRP	CD2	-38.48	-29.53	57.12	13.57
178	TRP	CE2	-38.10	-30.61	56.30	13.92
178	TRP	CE3	-38.73	-29.77	58.48	15.49
178	TRP	CD1	-38.19	-28.77	55.04	12.10
178	TRP	NE1	-37.93	-30.12	55.03	13.58
178	TRP	CZ2	-37.95	-31.92	56.79	14.49
178	TRP	CZ3	-38.58	-31.07	58.97	16.39
178	TRP	CH2	-38.19	-32.13	58.12	16.81
178	TRP	С	-40.31	-25.13	57.51	10.66
178	TRP	0	-40.55	-24.19	56.75	11.76
179	ILE	N	-40.13	-24.95	58.82	11.11
179	ILE	CA	-40.11	-23.61	59.41	9.30
179	ILE	CB	-40.99	-23.50	60.66	11.02
179	ILE	CG2	-40.96	-22.10	61.21	3.09
179	ILE	CG1	-42.44	-23.88	60.33	9.87
179		CD1	-43.38	-23.79	61.52	10.20
179	ILE	С	-38.66	-23.46	59.79	10.91
179	ILE	0	-38.13	-24.27	60.55	13.14
180	ILE	N	-38.01	-22.45	59.23	12.47
180	ILE	CA	-36.60	-22.25	59.48	10.39
180	ILE	CB	-35.84	-22.35	58.14	4.50
180	ILE	CG2	-34.41	-21.90	58.27	4.54
180	ILE	CG1	-35.91	-23.78	57.61	6.63
180	ILE	CD1	-35.21	-24.85	58.48	10.08
180	ILE	С	-36.27	-20.95	60.18	12.35

180	ILE	0	-36.73	-19.88	59.76	18.36
181	LYS	N	-35.50	-21.05	61.27	9.02
181	LYS	CA	-35.06	-19.88	62.03	6.93
181	LYS	CB	-34.92	-20.19	63.52	8.20
181	LYS	CG	-34.55	-18.99	64.39	4.18
181	LYS	CD	-34.21	-19.41	65.80	5.00
181	LYS	CE	-33.83	-18.23	66.69	2.00
181	LYS	NZ	-33.38	-18.69	68.03	2.00
181	LYS	С	-33.71	-19.45	61.47	6.56
181	LYS	0	-32.76	-20.23	61.51	9.50
182	ASN	N	-33.63	-18.25	60.93	6.31
182	ASN	CA	-32.38	-17.79	60.36	7.51
182	ASN	CB	-32.62	-17.18	58.97	12.34
182	asn	CG	-31.37	-17.16	58.12	13.66
182	ASN	OD1	-30.38	-17.83	58.43	13.80
182	ASN	ND2	-31.40	-16.41	57.03	14.20
182	ASN	С	-31.67	-16.80	61.27	8.95
182	asn	0	-32.29	-16.18	62.12	9.35
183	SER	N	-30.37	-16.65	61.07	12.23
183	SER	CA	-29.55	-15.74	61.87	12.33
183	SER	CB	-28.41	-16.50	62.54	11.14
183	SER		-27.51	-17.06	61.60	2.83
183	SER	C	-29.03	-14.56	61.05	17.18
183	SER	0	-27.84	-14.23	61.12	14.11
184	TRP	N	-29.92	-13.92	60.30	20.03
184	TRP	CA	-29.59	-12.77	59.45	17.75
	TRP		-29.94	-13.03	57.99	17.44
	TRP		-28.96	-13.87	57.26	11.74
	TRP		-29.09	-14.40	55.93	10.49
184	TRP	CE2	-27.91	-15.11	55.65	11.82
184			-30.09	-14.34	54.96	7.76
	TRP		-27.75	-14.27	57.71	13.64
	TRP		-27.11	-15.01	56.76	18.22
184	TRP	CZ2	-27.70	-15.76	54.44	8.06
184	TRP	CZ3	-29.88	-14.99	53.75	8.86
184	TRP	CH2	-28.70	-15.69	53.50	6.44
184	TRP	С	-30.30	-11.51	59.91	18.70
184	TRP	0	-30.21	-10.47	59.27	18.61
185	GLY	N	-31.05	-11.62	61.01	19.92
185	GLY	CA		-10.48	61.55	17.99
185	GLY	C	-33.25		61.47	21.34
185	GLY	0	-33.78	-11.41	60.68	24.20

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186 GLU N	-33.94	-9.85	62.30	25.86
186 GLU CA	-35.39	-9.85	62.34	28.63
186 GLU CB	-35.92	-9.06	63.55	33.72
186 GLU CG	-37.35	-9.41	64.01	33.62
186 GLU CD	-37.39	-10.59	64.97	33.61
186 GLU OE1	-36.64	-10.57	65.96	35.89
186 GLU OE2	-38.18	-11.54	64.74	33.82
186 GLU C	-35.93	-9.26	61.04	27.44
186 GLU O	-37.08	-9.48	60.68	29.63
187 ASN N	-35.09	-8.50	60.33	28.18
187 ASN CA	-35.54	-7.88	59.08	30.53
187 ASN CB	-35.10	-6.41	58.99	33.02
187 ASN CG	-36.04	-5.57	58.12	36.45
187 ASN OD1	-35.65	-4.51	57.62	41.13
187 ASN ND2	-37.28	-6.03	57.95	34.46
187 ASN C	-35.20	-8.65	57.79	29.17
187 ASN O	-35.12	-8.08	56.69	26.38
188 TRP N	-34.95	-9.94°	57.93	26.78
188 TRP CA	-34.69	-10.76	56.76	22.32
188 TRP CB	-33.39	-11.54	56.84	20.90
188 TRP CG	-33.28	-12.47	55.69	18.19
188 TRP CD2	-33.78	-13.82	55.64	20.16
188 TRP CE2	-33.53	-14.30	54.34	16.93
188 TRP CE3	-34.43	-14.65	56.56	15.73
188 TRP CD1	-32.75	-12.21	54.46	12.09
188 TRP NE1	-32.90	-13.31	53.64	17.84
188 TRP CZ2	-33.90	-15.58	53.94	18.81
188 TRP CZ3	-34.79	-15.92	56.17	18.58
188 TRP CH2	-34.53	-16.38	54.87	19.87
188 TRP C	-35.87	-11.70	56.68	21.57
188 TRP O	-36.40	-12.11	57.72	19.48
189 GLY N	-36.28	-12.04	55.46	22.26
189 GLY CA	-37.41	-12.93	55.26	19.03
189 GLY C	-38.59	-12.53	56.12	16.83
189 GLY O	-38.70	-11.37	56.51	16.64
190 ASN N	-39.44	-13.48	56.47	19.64
190 ASN CA	-40.60	-13.17	57.30	21.04
190 ASN CB	-41.76	-14.13	57.02	18.69
190 ASN CG	-43.09	-13.58	57.49	20.80
190 ASN OD1	-43.14	-12.58	58.20	25.63
190 ASN ND2	-44.18	-14.23	57.09	16.08
190 ASN C	-40.20	-13.18	58.78	20.26

190 ASN O	-40.08		59.39	21.80
191 LYS N	-39.91		59.31	19.09
191 LYS CA	-39.51		60.71	17.89
191 LYS CB	-40.69		61.66	19.60
191 LYS CG	-41.88		61.39	20.04
191 LYS CD	-43.02		62.36	17.35
191 LYS CE	-44.34		61.84	16.88
191 LYS NZ	-45.47		62.77	17.42
191 LYS C	-38.33	-12.71	61.12	14.97
191 LYS 0	-38.28	-13.22	62.23	11.80
192 GLY N	-37.38	-12.87	60.20	16.54
192 GLY CA	-36.20	-13.67	60.47	10.79
192 GLY C	-36.40	-15.14	60.19	12.10
192 GLY O	-35.53	-15.94	60.52	14.29
193 TYR N	-37.53	-15.50	59.59	15.02
193 TYR CA	-37.82	-16.91	59.26	15.85
193 TYR CB	-39.02	-17.45	60.08	19.04
193 TYR CG	-38.75	-17.68	61.55	20.49
193 TYR CD1	-38.79	-16.64	62.45	23.39
193 TYR CE1	-38.52	-16.84	63.79	24.79
193 TYR CD2	-38.43	-18.95	62.02	21.17
193 TYR CE2	-38.15	-19.17	63.36	22.23
193 TYR CZ	-38.19	-18.11	64.24	24.79
193 TYR OH	-37.88	-18.29	65.56	25.50
193 TYR C	-38.11	-17.13	57.77	14.84
193 TYR O	-38.58	-16.22	57.07	11.51
194 ILE N	-37.90	-18.36	57.33	16.12
194 ILE CA	-38.16	-18.78	55.96	14.70
194 ILE CB	-36.84	-18.88	55.15	17.12
194 ILE CG2	-35.85	-19.82	55.81	13.78
194 ILE CG1	-37.11	-19.29	53.70	15.67
194 ILE CD1	-35.97	-18.96	52.77	13.44
194 ILE C	-38.92	-20.12	55.97	12.70
194 ILE O	-38.59	-21.02	56.73	11.58
195 LEU N	-39.99	-20.21	55.18	12.52
195 LEU CA	-40.79	-21.43	55.09	12.23
195 LEU CB	-42.27	-21.12	54.93	12.99
195 LEU CG	-42.88	-20.23	55.99	16.19
195 LEU CD1	-44.36	-20.05		16.70
195 LEU CD2	-42.69	-20.87		17.95
195 LEU C	-40.36	-22.30	53.93	10.25
195 LEU O	-41.15	-22.59	53.05	15.85

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196 MET N	-39.12	2 -22.76	53.96	9.10
196 MET CA			-0.50	
196 MET CB	-37.15			
196 MET CG	-36.23			
196 MET SD	-34.62	-23.28		_
196 MET CE	-33.88	-23.71		8.14
196 MET C	-39.43			
196 MET O	-40.21			— <del>-</del>
197 ALA N	-39.26	-25.28	51.30	13.91
197 ALA CA	-40.03	-26.42	50.82	11.72
197 ALA CB	-39.92	-26.52	49.30	12.43 7.27
197 ALA C	-39.62			
197 ALA O	-38.45	-27.92		11.55
198 ARG N	-40.58	-28.64		11.95
198 ARG CA	-40.38		52.30	13.06
198 ARG CB	-41.05		53.69	14.09
198 ARG CG	-41.05	-31.27	54.39	5.88
198 ARG CD	-41.44	-31.09	55.84	6.02
198 ARG NE	-42.82	-30.68	56.02	3.48
198 ARG CZ	-43.85	-31.52	56.05	6.84
198 ARG NH1	-43.64	-32.82	55.89	2.00
198 ARG NH2	~45.07	-31.08	56.35	2.52
198 ARG C	-40.94	-31.02		12.96
198 ARG 0	-41.97	-30.84	50.78	15.58
199 ASN N	-40.29	-32.18	51.47	12.90
199 ASN CA		-33.34	50.66	13.45
199 ASN CB		-33.57	50.63	17.26
199 ASN CG	-42.69	-34.17	51.91	22.47
199 ASN OD1	-42.08	-35.09	52.47	20.71
199 ASN ND2	-43.83	-33.68	52.36	26.83
199 ASN C	-40.12	-33.20	49.24	12.34
199 ASN 0	-39.63	-34.18	48.69	12.85
200 LYS N	-40.22	-32.00	48.68	10.89
200 LYS CA		-31.64	47.34	8.22
200 LYS CB	-39.86	-30.13	47.11	8.55
200 LYS CG	-41.26	-29.57	47.04	5.90
200 LYS CD		-30.04	45.79	9.80
200 LYS CE		-29.09	45.40	9.34
200 LYS NZ		-27.88	44.72	10.27
200 LYS C		-32.09	47.02	7.56
200 LYS 0		-31.27	46.71	5.62
201 ASN N	-38.05	-33.39	47.07	9.28

201 ASN CA	-36.76	-33.99	46.79	10.44
201 ASN CB	-36.58	-34.11	45.28	12.24
201 ASN CG	-37.80	-34.72	44.61	16.00
201 ASN OD1	-38.18	-35.86		
201 ASN ND2	-38.47	-33.93		
201 ASN C	~35.55	-33.38		
201 ASN 0	-34.60	-32.95		
202 ASN N	-35.59	-33.36		
202 ASN CA	-34.50	-32.83	49.61	9.85
202 ASN CB	-33.31	-33.79	49.55	13.31
202 ASN CG	-32.38	-33.63	50.72	13.23
202 ASN OD1	-32.82	-33.40	51.84	16.24
202 ASN ND2	-31.09	-33.81	50.47	7.73
202 ASN C	-34.09	-31.43	49.19	
202 ASN O	-32.90	-31.12	49.17	13.84
203 ALA N	-35.05	-30.59	48.88	15.00
203 ALA H	-35.55	-31.01	49.01	15.00
203 ALA CA	-34.83		48.41	15.00
203 ALA CB	-35.54		48.10	15.00
203 ALA C	-33.66		49.16	15.00
203 ALA O	-33.48	-28.72	50.36	18.81
204 CYS N	-32.82	-27.85	48.40	16.60
204 CYS CA	-31.68	-27.12	48.94	13.11
204 CYS C	-30.69	-27.92	49.78	13.70
204 CYS O	-29.77		50.36	13.59
204 CYS CB	-32.14	-25.88	49.70	11.09
204 CYS SG	-32.81	-24.55	48.65	21.00
205 GLY N	-30.86	-29.23	49.82	15.46
205 GLY CA	-29.96	-30.07	50.60	17.31
205 GLY C	-30.16		52.09	18.03
205 GLY 0		-29.94	52.87	16.28
206 ILE N	-31.42	-29.72	52.49	18.33
206 ILE CA	-31.76	-29.48	53.88	18.94
206 ILE CB	-33.23	-28.93	54.04	19.06
206 ILE CG2		-29.85	53.34	18.93
206 ILE CG1	-33.59	-28.78	55.52	14.65
206 ILE CD1	-34.76	-27.88	55.75	16.36
206 ILE C	-31.52	-30.66	54.82	21.37
206 ILE 0	-31.03	-30.47	55.94	21.08
207 ALA N	-31.79	-31.88	54.34	20.23
207 ALA CA		-33.08	55.16	19.58
207 ALA CB	-32.79	-34.05	54.92	20.40

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207 ALA C	-30.20			
207 ALA O	-30.29			
208 ASN N	-30.18			
208 ASN CA	-29.26			17.17
208 ASN CA	-27.97	· -		16.44
208 ASN CB	-27.26			16.56
208 ASN CG 208 ASN OD1	-27.95			12.14
	-28.79			6.67
208 ASN ND2	-27.58		50.72	12.22
208 ASN C	-27.01		55.59	16.20
208 ASN O	-26.14			19.88
209 LEU N	-27.14		56.30	11.98
209 LEU CA	-26.22		57.40	12.42
209 LEU CB	-25.19		56.96	11.33
209 LEU CG	-23.81	-31.03	57.61	6.60
209 LEU CD1	-23.24	-32.42	57.81	6.26
209 LEU CD2	-22.89	-30.23	56.73	6.70
209 LEU C	-26.87	-31.77	58.73	14.26
209 LEU O	-26.32		59.48	15.18
210 ALA N	-27.99	-32.44	59.08	15.00
210 ALA H		-32.84	58.30	15.00
210 ALA CA	-28.76	-32.10	60.27	15.00
210 ALA CB	-30.25	-32.37	60.06	15.00
210 ALA C	-28.31	-32.97	61.45	15.00
210 ALA O	-28.07	-34.16	61.40	11.82
211 SER N	-28.32	-32.32	62.66	12.69
211 SER CA	-27.97	-33.00	63.90	11.35
211 SER CB	-26.47	-33.22	64.03	10.79
211 SER OG	-25.80	-32.01	64.32	5.14
211 SER C	-28.50	-32.25	65.11	11.89
211 SER O	-28.51	-31.01	65.14	11.91
212 PHE N	-28.97	-32.99	66.09	12.57
212 PHE CA	-29.47	-32.38	67.31	11.64
212 PHE CB	-30.95	-32.69	67.51	12.77
212 PHE CG		-34.12	67.44	11.84
212 PHE CD1	-31.61	-34.70	66.24	10.42
212 PHE CD2		-34.91	68.58	14.45
212 PHE CE1		-36.03	66.17	16.06
212 PHE CE2		-36.24	68.53	15.41
212 PHE CZ		-36.81	67.32	16.55
212 PHE C		-32.83	68.48	12.10
212 PHE O		-33.85	68.41	12.10
213 PRO N		-32.05	69.57	12.03
		· <b>-</b> -		14.73

213 PRO CD	-29.24	-30.73	69.77	10.69
213 PRO CA	-27.79	-32.41	70.74	12.29
213 PRO CB	-27.41	-31.05	71.29	11.67
213 PRO CG	-28.70	-30.29	71.12	
213 PRO C	-28.56	-33.23		
213 PRO O	-29.78	-33.11		
214 LYS N	-27.82	-34.09	72.45	14.45
214 LYS CA	-28.39	-34.94	73.50	
214 LYS CB	-27.80	-36.34	73.46	
214 LYS CG	-28.06	-37.08	72.17	24.16
214 LYS CD	-27.54		72.24	29.93
214 LYS CE	-26.02	-38.57	72.40	33.14
214 LYS NZ	-25.49	-39.97	72.44	35.37
214 LYS C	-28.15	-34.26	74.84	21.45
214 LYS O	-27.02		75.20	16.74
215 MET N	-29.23	-34.05	75.57	22.65
215 MET CA	-29.16	-33.38	76.85	25.38
215 MET CB	-30.04	-32.13	76.79	27.81
215 MET CG	-29.77	-31.09	77.85	26.22
215 MET SD	-30.44	-29.54	77.29	25.84
215 MET CE	-28.92	-28.76	76.63	23.17
215 MET C	-29.56	-34.30	77.99	25.33
215 MET OT1	-28.65	-34.77	78.71	27.00
215 MET OT2	-30.77	-34.58	78.12	23.48
216 нон он2	-31.11	-16.42	65.02	14.43
217 нон он2	-29.30	-20.25	62.17	18.73
218 нон он2	-10.67	-12.22	63.70	43.10
219 нон он2	-16.45	-12.20	72.96	5.87
220 НОН ОН2	-35.12	-23.55	69.64	9.44
221 нон он2	-24.01	-30.97	61.16	4.73
222 НОН ОН2	-13.01	-8.39	61.94	32.49
223 нон он2	-14.66	-21.66	66.41	2.00
224 НОН ОН2	-43.65	-26.52	48.92	29.00
225 нон он2		-35.43	55.56	13.84
226 нон он2	-40.92		68.90	12.03
227 НОН ОН2		-25.26	44.65	38.82
228 НОН ОН2	-24.12	-5.83	68.94	43.50
229 НОН ОН2	-30.71	-18.60	67.86	32.89
230 нон он2		-26.71	51.39	30.11
231 нон он2		-24.80	49.94	8.69
232 нон он2	-46.77	-33.25	57.36	12.67
233 нон он2	-28.91	-10.19	75.44	15.32

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234 НОН ОН2	-36.31	-14.76	75.60	16.14
235 нон он2	~16.18		68.62	27.92
236 НОН ОН2	-16.52	-8.98	75.02	28.63
237 НОН ОН2	-10.50	-18.37		39.29
238 НОН ОН2	-9.29			29.36
239 НОН ОН2	-45.95		54.42	32.47
240 HOH OH2	-33.98		44.88	32.74
241 HOH OH2	-36.55		52.42	11.87
242 HOH OH2	-41.73	-34.84	55.47	18.73
243 нон он2	-41.21	-19.79	71.20	12.36
244 нон он2	-47.90		72.76	
245 HOH OH2	-42.20	-14.92	70.78	27.97
246 HOH OH2	-26.14	-8.98	67.92	34.65 37.03
247 HOH OH2	-32.81	-7.84	63.95	
248 нон он2	-19.95	-7.54	63.08	35.20 32.69
249 нон он2	-16.19	-10.67	61.92	30.22
250 нон он2	-35.01	-39.74	65.90	10.75
251 нон он2	-13.63	-24.17	76.67	16.76
252 нон он2	-8.21	-25.68	60.56	19.27
253 нон он2	-20.14	-27.62	51.69	32.97
254 НОН ОН2	-25.01	-33.27	60.23	25.95
255 нон он2	-13.56	-29.77	72.49	36.07
256 нон он2	-33.43	-40.03	63.75	25.23
257 нон он2	-20.84	-23.81	87.06	28.06
258 нон он2	-13.50	-12.46	62.97	41.75
259 нон он2	-28.41	-30.03	56.55	11.99
260 нон он2	-28.87	-16.56	42.17	23.76
261 HOH OH2	-25.56	-19.23	42.44	8.60
262 НОН ОН2	-32.08	-33.98	47.05	47.25
263 НОН ОН2	-22.35	-31.23	43.22	18.34
264 НОН ОН2	-32.62	-28.89	42.32	42.83
265 нон он2		-30.05	50.12	38.26

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (A<sup>2</sup>) for the cathepsin K complex with inhibitor 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone.

Residue Atom	X	Y	z	В
1 ALA CB	-46.30	-38.07	64.13	15.00
1 ALA C	-48.64	-37.16	64.46	15.00
1 ALA O	-49.53	-37.68	63.79	15.00
1 ALA N	-48.10	-39.41	65.31	15.00
1 ALA CA	-47.55	-38.03	65.08	15.00
2 PRO N	-48.69	-35.88	64.81	15.00
2 PRO CD	-48.06	-35.33	66.02	15.00
2 PRO CA	-49.67	-34.93	64.29	15.00
2 PRO CB	-49.53	-33.73	65.24	15.00
2 PRO CG	-48.13	-33.85	65.77	15.00
2 PRO C	-49.42	-34.53	62.86	15.00
2 PRO O	-48.34	-34.77	62.28	15.00
3 ASP N	-50.40	-33.85	62.30	15.00
3 ASP CA	-50.29	-33.37	60.93	15.00
3 ASP CB	-51.65	-33.42	60.27	15.00
3 ASP CG	-52.12	-34.84	60.00	15.00
3 ASP OD1	-52.02	-35.30	58.84	15.00
3 ASP OD2	-52.59	-35.51	60.95	15.00
3 ASP C	-49.76	-31.93	61.01	15.00
3 ASP O	-49.44	-31.32	59.99	15.00
4 SER N	-49.58	-31.45	62.24	15.00
4 SER CA	-49.13	-30.09	62.51	15.00
4 SER CB	-50.32	-29.14	62.52	15.00
4 SER OG	-50.86	-29.02	61.20	15.00
4 SER C	-48.39	-29.96	63.84	15.00
4 SER O	-48.74	-30.63	64.84	15.00
5 VAL N	-47.34	-29.15	63.84	15.00
5 VAL CA	-46.50	-28.91	64.99	15.00
5 VAL CB	-45.35	-29.97	65.11	15.00
5 VAL CG1	-44.17	-29.43	65.89	15.00
5 VAL CG2	-45.84	-31.21	65.74	15.00
5 VAL C	-45.86	-27.57	64.80	15.00
5 VAL O	-45.37	-27.25	63.70	15.00

6 ASP N	-45.84		65.89	15.00
6 ASP CA				
6 ASP CB	-46.16		65.46	
6 ASP CG	-45.40		65.15	
6 ASP OD1	-44.21		65.51	15.00
6 ASP OD2	-45.98		64.51	15.00
6 ASP C	-44.81		67.34	
6 ASP O	-45.65		68.17	15.00
7 TYR N	-43.51		67.61	15.00
7 TYR CA	-43.00		68.94	15.00
7 TYR CB	-41.63		69.13	15.00
7 TYR CG	-41.68	-27.30	69.31	15.00
7 TYR CD1		-27.88	70.57	15.00
7 TYR CE1	-41.60	-29.29	70.74	15.00
7 TYR CD2	-41.86	-28.14	68.22	15.00
7 TYR CE2	-41.91	-29.55	68.38	15.00
7 TYR CZ	-41.79	-30.10	69.64	15.00
7 TYR OH	-41.91	-31.46	69.83	15.00
7 TYR C	-42.85	-23.70	69.30	15.00
7 TYR O	-42.16	-23.39	70.26	15.00
8 ARG N	-43.40	-22.80	68.50	15.00
8 ARG CA	-43.31	-21.39	68.80	15.00
8 ARG CB	-43.55	-20.55	67.56	15.00
8 ARG CG	-42.37	-20.55	66.60	15.00
8 ARG CD	-42.57	-19.68	65.38	15.00
8 ARG NE	-43.64	-20.20	64.53	15.00
8 ARG CZ	-43.88	-19.78	63.29	15.00
8 ARG NH1	-43.09	-18.84	62.77	15.00
8 ARG NH2	-44.92	-20.26	62.60	15.00
8 ARG C	-44.37	-21.14	69.84	15.00
8 ARG O	-44.09	-20.64	70.93	15.00
9 LYS N	-45.58	-21.65	69.54	15.00
9 LYS CA	<b>-46</b> .73	-21.55	70.42	15.00
9 LYS CB	-47.95	-22.19	69.75	15.00
9 LYS CG		-21.37	68.55	15.00
9 LYS CD	-48.71	-22.24	67.29	15.00
9 LYS CE	-49.41	-21.46	66.17	15.00
9 LYS NZ		21.36	66.45	15.00
9 LYS C		-22.17	71.79	15.00
9 LYS O	-46.36	-21.45	72.77	15.00
10 LYS N		-23.46	71.83	15.00
10 LYS CA	-45.85	-24.13	73.10	15.00
10 LYS CB	-45.43	-25.57	72.86	15.00

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10 LYS CG	-46.52	-26.39	72.25	15.00
10 LYS CD	-46.05	-27.76		
10 LYS CE	-46.89	-28.35	70.81	15.00
10 LYS NZ	-48.31	-27.94	70.92	15.00
10 LYS C	-44.81	-23.41	73.94	15.00
10 LYS O	-44.74	-23.62	75.14	
11 GLY N	-43.95	-22.63	73.30	15.00
11 GLY CA	-42.94	-21.90	74.05	15.00
11 GLY C	-41.53	-22.43	74.01	15.00
11 GLY O	-40.78	-22.24	74.97	15.00
12 TYR N	-41.13	-23.07	72.91	15.00
12 TYR CA	-39.76	-23.60	72.84	15.00
12 TYR CB	-39.78	-24.98	72.15	15.00
12 TYR CG	-40.42	-26.09	72.97	15.00
12 TYR CD1	-41.76	-26.06	73.35	15.00
12 TYR CE1	-42.34	-27.10	74.13	15.00
12 TYR CD2	-39.66	-27.18	73.39	15.00
12 TYR CE2	-40.21	-28.21	74.17	15.00
12 TYR CZ	-41.55	-28.17	74.53	15.00
12 TYR OH	-42.10	-29.21	75.25	15.00
12 TYR C	-38.82	-22.68	72.08	15.00
12 TYR O	-37.64	-22.97	71.90	15.00
13 VAL N	-39.37	-21.59	71.56	15.00
13 VAL CA	-38.56	-20.71	70.75	15.00
13 VAL CB	-39.14	-20.70	69.33	15.00
13 VAL CG1	-38.15	-20.04	68.35	15.00
13 VAL CG2	-39.49	-22.09	68.92	15.00
13 VAL C	-38.31	-19.27	71.23	15.00
13 VAL 0	-39.24	-18.52	71.45	15.00
14 THR N	-37.03	-18.90	71.32	15.00
14 THR CA	-36.6 <u>4</u>	-17.56	71.76	15.00
14 THR CB	-35.17	-17.57	72.25	15.00
14 THR OG1	-34.33	-18.03	71.18	15.00
14 THR CG2	-35.01	-18.43	73.50	15.00
14 THR C	-36.70	-16.58	70.60	15.00
14 THR O	-37.20	-16.92	69.53	15.00
15 PRO N	-36.27	-15.32	70.83	15.00
15 PRO CD	-36.21	-14.64	72.14	15.00
15 PRO CA	-36.29	-14.32	69.76	15.00
L5 PRO CB	-35.98	-13.04	70.49	15.00
L5 PRO CG	-36.64	-13.24	71.83	15.00
15 PRO C	-35.23	-14.60	68.72	15.00
.5 PRO O	-34.20	-15.21	69.02	15.00

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16 VAL N	-35.48	-14.05	67.52	15.00
16 VAL CA	-34.61	-14.19		15.00
16 VAL CB	-35.37	-13.74	65.10	15.00
16 VAL CG1	-34.43	-13.61	63.92	15.00
16 VAL CG2	-36.44	-14.72	64.78	15.00
16 VAL C	-33.40	-13.33	66.56	15.00
16 VAL O	-33.47			
17 LYS N	-32.24	-13.78		
17 LYS CA	-31.04			15.00
17 LYS CB	-29.96	-13.71	67.06	
17 LYS CG	-30.45	-14.44	68.35	15.00
17 LYS CD	-30.63	-13.52	69.54	15.00
17 LYS CE	-30.89	-14.30	70.82	15.00
17 LYS NZ	-32.14	-15.10	70.83	15.00
17 LYS C	-30.46	-12.58	64.91	15.00
17 LYS O	-31.06		63.87	15.00
18 ASN N	-29.30	-11.94	64.94	15.00
18 ASN CA	-28.58	-11.48	63.75	15.00
18 ASN CB	-28.58	-9.96	63.69	15.00
18 ASN CG	-29.93	-9.39	63.96	15.00
18 ASN OD1	-30.89	-9.75	63.32	15.00
18 ASN ND2	-30.02	-8.56	64.97	15.00
18 ASN C	-27.16	-11.93	63.99	15.00
18 ASN O	-26.58	-11.59	65.03	15.00
19 GLN N	-26.59	-12.69	63.06	15.00
19 GLN CA	-25.22	-13.13	63.22	15.00
19 GLN CB	-24.96	-14.41	62.43	15.00
19 GLN CG	-25.34	-14.25	60.98	15.00
19 GLN CD	-25.10		60.16	15.00
19 GLN OE1	-26.04		59.82	15.00
19 GLN NE2	-23.86	-15.72	59.77	15.00
19 GLN C	-24.24	-12.02	62.83	15.00
19 GLN O	-23.05	-12.15	63.08	15.00
20 GLY N	-24.72	-11.01	62.11	15.00
20 GLY CA	-23.86	-9.91	61.69	15.00
20 GLY C	-22.83	-10.31	60.65	15.00
20 GLY O	-23.08	-11.26	59.94	15.00
21 GLN N	-21.69	-9.61	60.49	15.00
21 GLN CA	-20.72	-10.02	59.43	15.00
21 GLN CB	-20.01	-8.83	58.75	15.00
21 GLN CG	-20.87	-8.13	57.66	15.00
21 GLN CD	-21.58	-9.06	56.63	J.5 . 00
21 GLN OE1	-20.99	-10.02	56.16	15.00

21 GLN NE2	-22.84		56.29	15.00
21 GLN C	-19.72		59.87	15.00
21 GLN O	-18.50		59.66	15.00
22 CYS N	-20.32		60.26	15.00
22 CYS CA	-19.59	-13.37	60.72	15.00
22 CYS C	-20.26		60.30	15.00
22 CYS O	-21.48		60.28	
22 CYS CB	-19.48		62.25	15.00
22 CYS SG	-19.02		63.24	15.00
23 GLY N	-19.45	-15.65	59.90	15.00
23 GLY CA	-20.01	-16.93	59.52	15.00
23 GLY C	-20.27	-17.81	60.75	15.00
23 GLY O	-19.73	-18.91	60.83	15.00
24 SER N	-21.20	-17.40	61.61	15.00
24 SER CA	-21.49	-18.12	62.84	15.00
24 SER CB	-21.32	-17.21	64.08	15.00
24 SER OG	-22.22	-16.10	64.07	15.00
24 SER C	-22.89	-18.73	62.86	15.00
24 SER O	-23.44	-19.03	63.90	15.00
25 CYS N	-23.46	-18.96	61.69	15.00
25 CYS CA	-24.78	-19.55	61.56	15.00
25 CYS CB	-25.18	-19.59	60.07	15.00
25 CYS SG	-24.19	-20.73	59.02	15.00
25 CYS C	-24.80	-20.92	62.24	15.00
25 CYS O	-25.77	-21.25	62.88	15.00
25 INH C1	-14.75	-27.52	59.81	15.00
25 INH C2	-15.58	-26.77	58.94	15.00
25 INH C3	-15.24	-25.44	58.65	15.00
25 INH C4	-14.06	-24.88	59.25	15.00
25 INH C5	-13.21	-25.64	60.14	15.00
25 INH C6	-13.57	-26.96	60.42	15.00
25 INH C7	-16.11	-24.63	57.72	15.00
25 INH 08	-17.39	-25.29	57.48	15.00
25 INH .C9	-18.43	-24.53	57.00	15.00
25 INH 010	-18.33	-23.63	56.17	15.00
25 INH N11	-19.57	-24.86	57.54	15.00
25 INH C12	-20.88	-24.22	57.23	15.00
25 INH C13	-21.31	-23.29	58.42	15.00
25 INH N14	-21.06	-21.86	58.16	15.00
25 INH C15	-21.68	-21.41	56.87	15.00
25 INH C16	-21.59	-21.00	59.27	15.00
25 INH C17	-22.57	-20.07	58.55	15.00
25 INH C18	-22.15	-19.99	57.10	15.00

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25 INH C19			57.01	15.00
25 INH C20	_	-25.74		
25 INH C21			58.28	
25 INH C22		. –		
25 INH N23			56.18	
25 INH C24	_		55.84	
25 INH 025				
25 INH C26	=		54.91	
25 INH C27			53.43	
25 INH C28	-25.22			15.00
25 INH C29			51.66	15.00
25 INH C30	-25.25		51.69	15.00
25 INH N31	-24.30	-16.10	54.92	15.00
25 INH C32	-24.85	-15.34	55.87	15.00
25 INH 033	-25.40			15.00
25 INH 034		-13.96	55.64	15.00
25 INH C35	· · · · <del>·</del>		56.80	15.00
25 INH C36		-11.91	56.62	15.00
25 INH C37	-25.21	-10.60	56.86	15.00
25 INH C38	-26.09			15.00
25 INH N39		-9.73	56.40	15.00
25 INH C40		-10.96	56.16	15.00
25 INH C41	·	-12.08	56.26	15.00
25 INH 042	-22.66	-18.72	58.88	15.00
26 TRP N	-23.72	-21.69	62.15	15.00
26 TRP CA	-23.64	-23.01	-	15.00
26 TRP CB	-22.30	-23.65		15.00
26 TRP CG		-22.75	63.06	15.00
26 TRP CD2	-20.40	-22.85	64.28	15.00
26 TRP CE2		-21.79	64.22	15.00
26 TRP CE3	-20.41	-23.71	65.38	15.00
26 TRP CD1	-20.71	-21.70		15.00
26 TRP NE1 26 TRP CZ2	-19.66	-21.12		15.00
	-18.49		65.24	15.00
26 TRP CZ3	-19.48		66.41	15.00
26 TRP CH2		-22.41	66.33	15.00
26 TRP C		-22.83	64.40	15.00
26 TRP 0		-23.79	65.16	15.00
27 ALA N		-21.54	64.91	15.00
27 ALA CA		-21.43	66.36	15.00
27 ALA CB			66.80	15.00
27 ALA C		-21.21		15.00
27 ALA O	-25.52	-21.89	67.60	15.00

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28 PHE N	-25.63	-20.27	66.03	15.00
28 PHE CA	-27.02	-19.90	66.31	15.00
28 PHE CB	-27.46	-18.71	65.44	15.00
28 PHE CG	-26.88		65.85	15.00
28 PHE CD1	-25.73		65.27	15.00
28 PHE CD2	-27.47	-16.73	66.88	15.00
28 PHE CE1	-25.16		65.70	15.00
28 PHE CE2	-26.92		67.33	15.00
28 PHE CZ	-25.75		66.75	15.00
28 PHE C	-27.93		66.06	15.00
28 PHE O	-29.08		66.44	15.00
29 SER N	-27.44	-22.09	65.33	15.00
29 SER CA	-28.24	-23.28	65.05	15.00
29 SER CB	-27.74	-23.99	63.81	15.00
29 SER OG	-28.44		63.58	15.00
29 SER C	-28.13	-24.15	66.27	15.00
29 SER 0	-29.14	-24.63	66.79	15.00
30 SER N	-26.90	-24.31	66.76	15.00
30 SER CA	-26.64	-25.13	67.94	15.00
30 SER CB	-25.14	-25.24	68.21	15.00
30 SER OG	-24.40	-25.67	67.06	15.00
30 SER C	-27.29	-24.63	69.21	15.00
30 SER O	-27.66	-25.40	70.08	15.00
31 VAL N	-27.31	-23.32	69.35	15.00
31 VAL CA	-27.89	-22.67	70.51	15.00
31 VAL CB	-27.39	-21.19	70.58	15.00
31 VAL CG1	-28.13	-20.34	71.59	15.00
31 VAL CG2	-25.91	-21.19	70.86	15.00
31 VAL C	-29.41	-22.80	70.55	15.00
31 VAL O	-30.02	-22.96	71.63	15.00
32 GLY N	-29.99	-22.81	69.35	15.00
32 GLY CA	-31.43	-22.91	69.21	15.00
32 GLY C	-31.90	-24.26	69.66	15.00
32 GLY O	-32.81		70.47	15.00
33 ALA N	-31.22		69.17	15.00
33 ALA CA	-31.51	-26.66	69.53	15.00
33 ALA CB	-30.58	-27.55	68.80	15.00
33 ALA C	-31.38	-26.86	71.06	15.00
33 ALA O	-32.23	-27.54	71.68	15.00
34 LEU N	-30.29		71.63	15.00
34 LEU CA		-26.45	73.05	15.00
34 LEU CB		-25.82	73.41	15.00
34 LEU CG	-27.43	-26.58	73.17	15.00

		TABLE !	1111	
34 LEU CD1	-26.28		73.53	15.00
34 LEU CD2	-27.38		74.05	15.00
34 LEU C	-31.12	-25.73	73.83	15.00
34 LEU O	-31.65	-26.29	74.80	15.00
35 GLU N	-31.47	-24.51	73.44	15.00
35 GLU CA	-32.54	-23.76	74.12	15.00
35 GLU CB	-32.78	-22.42	73.41	15.00
35 GLU CG	-31.67	-21.39	73.64	15.00
35 GLU CD	-31.63	-20.29	72.57	15.00
35 GLU OE1	-32.41	-20.37	71.57	15.00
35 GLU OE2	-30.81	-19.35	72.71	15.00
35 GLU C	-33.83	-24.57	74.20	15.00
35 GLU O	-34.34	-24.83	75.31	15.00
36 GLY N	-34.26	-25.10	73.05	15.00
36 GLY CA	-35.48	-25.91	72.99	15.00
36 GLY C	-35.58	-27.11	73.93	15.00
36 GLY O	-36.67	-27.47	74.37	15.00
37 GLN N	-34.44	-27.72	74.24	15.00
37 GLN CA	-34.40	-28.87	75.12	15.00
37 GLN CB	-33.06	-29.60	74.95	15.00
37 GLN CG	-32.79	-30.19	73.58	15.00
37 GLN CD	-33.83	-31.19	73.19	15.00
37 GLN OE1	-33.99	-32.23	73.84	15.00
37 GLN NE2	-34.56	-30.89	72.11	15.00
37 GLN C	-34.52	-28.38	76.55	15.00
37 GLN O	-35.01	-29.10	77.43	15.00
38 LEU N	-33.98	-27.18	76.79	15.00
38 LEU CA	-34.03	-26.60	78.12	15.00
38 LEU CB	-33.30	-25.26	78.14	15.00
38 LEU CG	-33.24	-24.54	79.47	15.00
38 LEU CD1	-32.61	-25.51	80.46	15.00
38 LEU CD2	-32.41	~23.30	79.36	15.00
38 LEU C	-35.46	-26.45	78.60	15.00
38 LEU O	-35.86	-27.10	79.58	15.00
39 LYS N	-36.26	-25.70	77.84	15.00
39 LYS CA	-37.68	-25.45	78.15	15.00
39 LYS CB	-38.34	-24.71	76.97	15.00
39 LYS CG	-39.49	-23.74	77.33	15.00
39 LYS CD	-40.74	-24.43	77.83	15.00
39 LYS CE	-41.85	-23.45	78.12	15.00
39 LYS NZ	-41.65	-22.78	79.42	15.00
39 LYS C	-38.42	-26.78	78.46	15.00
39 LYS O	-39.33	-26.82	79.30	15.00

40 LYS N	-38.04	-27.84	77.77	15.00
40 LYS CA	-38.66	-29.14		15.00
40 LYS CB	-38.41	-30.07	76.81	15.00
40 LYS CG	-39.57	-31.02	76.53	15.00
40 LYS CD	-39.10	-32.29	75.82	15.00
40 LYS CE	-40.02	-32.72	74.68	15.00
40 LYS NZ	-40.21	-31.65		15.00
40 LYS C	-38.08	-29.74	79.27	15.00
40 LYS O	-38.79	-30.27	80.12	15.00
41 LYS N	-36.77	-29.60	79.41	15.00
41 LYS CA	-36.05	-30.13	80.55	15.00
41 LYS CB	-34.53	-29.93	80.33	15.00
41 LYS CG	-33.63	-30.86	81.11	15.00
41 LYS CD	-33.79	-32.26	80.55	15.00
41 LYS CE	-33.13	-33.31	81.42	15.00
41 LYS NZ	-33.45	-34.68	80.86	15.00
41 LYS C	-36.49		81.84	15.00
41 LYS O	-37.34		82.61	15.00
42 THR N	-35.96	-28.25	82.04	15.00
42 THR CA	-36.22	-27.48	83.24	15.00
42 THR CB	-34.87		83.78	15.00
42 THR OG1	-34.35	-25.91	82.92	15.00
42 THR CG2	-33.86	-28.07	83.79	15.00
42 THR C	-37.16	-26.30	83.02	15.00
42 THR O	-36.86	-25.20	83.45	15.00
43 GLY N	-38.28	-26.52	82.34	15.00
43 GLY CA	-39.26	-25.47	82.12	15.00
43 GLY C	-38.87	-24.03	81.79	15.00
43 GLY O	-39.77	-23.21	81.54	15.00
44 LYS N	-37.56	-23.75	81.75	15.00
44 LYS CA	-37.02	-22.41	81.49	15.00
44 LYS CB	-35.84	-22.13	82.44	15.00
44 LYS CG	-36.25	-21.41	83.72	15.00
44 LYS CD	-35.09	-21.25	84.67	15.00
44 LYS CE		-20.36	85.86	15.00
44 LYS NZ		-20.99	86.66	15.00
4 LYS C	-36.56	-22.13	80.06	15.00
14 LYS O	-35.71	-22.83	79.51	15.00
15 LEU N	-37.07	-21.04	79.50	15.00
15 LEU CA	-36.70	-20.66	78.14	15.00
5 LEU CB		-20.08	77.46	15.00
5 LEU CG	-38.06		75.94	15.00
5 LEU CD1	-37.93	-21.25	75.26	15.00

45 LEU CD2	-39.40	-19.29	75.63	15.00
45 LEU C	-35.66			15.00
45 LEU O	-36.00	-18.46	78.71	15.00
46 LEU N	-34.40	-19.94	78.29	
46 LEU CA	-33.30	-18.99		
46 LEU CB	-32.23	-19.62		
46 LEU CG	-31.90	-18.84	80.70	
46 LEU CD1	-32.48	-19.56		
46 LEU CD2	-30.35	-18.72	80.89	
46 LEU C	-32.68	-18.70	77.13	15.00
46 LEU O	-32.93	-19.45	76.19	15.00
47 ASN N		-17.61	76.99	15.00
47 ASN CA	-31.23		75.73	15.00
47 ASN CB	-31.29	-15.79	75.44	15.00
47 ASN CG	-32.61	-15.36	74.77	
47 ASN OD1	-32.68		73.55	15.00
47 ASN ND2	-33.63		75.58	15.00
47 ASN C	-29.74	-17.70	75.90	15.00
47 ASN 0	-29.01	-16.99	76.58	15.00
48 LEU N	-29.28	-18.80	75.29	15.00
48 LEU CA	-27.88	-19.20	75.45	15.00
48 LEU CB	-27.67	-20.68	75.16	15.00
48 LEU CG	-28.05	-21.69	76.27	15.00
48 LEU CD1	-27.81	-21.07	77.65	15.00
48 LEU CD2	-29.49	-22.12	76.12	15.00
48 LEU C	-26.85	-18.34	74.73	15.00
48 LEU O	-27.20	-17.31	74.23	15.00
49 SER N	-25.58	-18.70	74.74	15.00
49 SER CA	-24.59	-17.83	74.11	15.00
49 SER CB	-23.51	-17.50	75.14	15.00
49 SER OG	-22.32	-17.09	74.51	15.00
49 SER C	-23.93	-18.37	72.87	15.00
49 SER O	-23.08	-19.27	72.98	15.00
50 PRO N	-24.29	-17.86	71.67	15.00
50 PRO CD	-25.31	-16.83	71.34	15.00
50 PRO CA	-23.66	-18.36	70.45	15.00
50 PRO CB	-24.47	-17.67	69.35	15.00
50 PRO CG	-24.96	-16.44	69.97	15.00
50 PRO C	-22.18	-17.94	70.44	15.00
50 PRO O	-21.36	-18.62	69.86	15.00
51 GLN N	-21.82	-16.87	71.13	15.00
51 GLN CA	-20.43	-16.41	71.17	15.00
51 GLN CB		-14.98	71.74	15.00

51 GLN CG	_		71.69	15.00
51 GLN CD	_			
51 GLN OE1			69.76	
51 GLN NE2	-17.54			
51 GLN C	-19.46	-17.34		
51 GLN O	-18.31	-17.50		
52 ASN N	-19.90	-17.99	73.02	
52 ASN CA	-19.02			
52 ASN CB	-19.79	-19.58		
52 ASN CG	-18.92	-20.56		
52 ASN OD1	-19.45	-21.25	76.60	
52 ASN ND2	-17.61	-20.58	75.49	
52 ASN C	-18.54	-19.96	72.80	
52 ASN O	-17.34	-20.22		
53 LEU N	-19.50	-20.50	72.05	15.00
53 LEU CA	-19.28	-21.52	71.00	15.00
53 LEU CB	-20.62	-21.86	70.33	15.00
53 LEU CG	-21.43	-23.10	70.69	
53 LEU CD1	-21.19			15.00
53 LEU CD2	-22.91	-22.80	70.38	15.00
53 LEU C	-18.28	-21.04	69.94	15.00
53 LEU O	-17.34	-21.76	69.60	15.00
54 VAL N	-18.48	-19.82	69.46	15.00
54 VAL CA	-17.63	-19.27		15.00
54 VAL CB	-18.01	-17.82		15.00
54 VAL CG1	-16.94	-17.17	67.25	15.00
54 VAL CG2	-19.32	-17.76	67.40	15.00
54 VAL C	-16.18	-19.32	68.84	15.00
54 VAL O	-15.36	-19.90		15.00
55 ASP N	-15.88	-18.75		15.00
55 ASP CA	-14.53	-18.65	70.58	15.00
55 ASP CB	-14.49	-17.73	71.81	15.00
55 ASP CG	-14.86	-16.30	71.52	
55 ASP OD1	-14.87	-15.89	70.35	15.00
55 ASP OD2	-15.13	-15.56	72.49	15.00
55 ASP C	-13.87	-19.93	71.08	15.00
55 ASP O	-12.65	-20.14	70.82	15.00
56 CYS N	-14.62	-20.75	71.81	15.00
56 CYS CA	-14.06	-21.94	72.42	15.00
56 CYS C		-23.23	71.62	15.00
56 CYS O		-24.15	71.85	15.00
56 CYS CB		-22.17	73.76	15.00
56 CYS SG		-20.81	74.99	15.60
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57 VAL N	-15.00		70.62	15.00
57 VAL CA	-15.09		69.88	
57 VAL CB	-16.51	-24.71	69.30	
57 VAL CG1	-16.66	-26.03	68.70	
57 VAL CG2	-17.56	-24.53	70.42	
57 VAL C	-13.92	-24.66	68.88	
57 VAL 0	-14.05	-24.47	67.68	
58 SER N	-12.79	-25.08	69.43	
58 SER CA	-11.51	-25.26		15.00
58 SER CB	-10.43	-25.78	69.66	15.00
58 SER OG	-9.18	-25.96	68.99	15.00
58 SER C	-11.53	-26.13	67.45	15.00
58 SER O	-10.66		66.58	15.00
59 GLU N	-12.54			15.00
59 GLU CA	-12.67			15.00
59 GLU CB	-13.47		66.60	15.00
59 GLU CG	-13.12		68.00	15.00
59 GLU CD	-14.02	-29.15	69.13	15.00
59 GLU OE1	-15.16	-29.68	69.29	15.00
59 GLU OE2	-13.56		69.84	15.00
59 GLU C	-13.30	-27.07	65.05	15.00
59 GLU O	-13.21		63.91	15.00
60 asn n	-13.91	-25.93	65.35	15.00
60 ASN CA	-14.50	-25.14	64.28	15.00
60 ASN CB	-15.92	-24.71	64.63	15.00
60 ASN CG	-16.96	-25.87	64.48	15.00
60 ASN OD1	-17.94	-25.95	65.23	15.00
60 ASN ND2	-16.75	-26.74	63.50	15.00
60 ASN C	-13.60	-23.94	63.89	15.00
60 ASN O	-12.43	-23.91	64.25	15.00
61 ASP N	-14.11	-23.06	63.02	15.00
61 ASP CA	-13.36	-21.90		15.00
61 ASP CB	-13.29	-21.96	61.02	15.00
61 ASP CG	-11.96	-22.46	60.52	15.00
61 ASP OD1		-23.42	61.13	15.00
61 ASP OD2		-21.91	59.50	15.00
61 ASP C		-20.55	62.98	15.00
61 ASP O		-19.49	62.62	15.00
62 GLY N		-20.59	63.77	15.00
62 GLY CA	_	-19.35	64.30	15.00
62 GLY C			63.21	15.00
62 GLY O			62.59	15.00
63 CYS N			62.93	15.00
				~J.00

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63 CYS CA	-16.40	-16.58	61.91	15.00
63 CYS C	-16.18	-17.07	60.50	15.00
63 CYS O	-16.75	-16.57	59.55	15.00
63 CYS CB	-16.13	-15.11	62.08	
63 CYS SG	-17.00		63.54	
64 GLY N	-15.46		60.41	15.00
64 GLY CA	-15.19	-18.79	59.11	15.00
64 GLY C	-16.12	-19.97	58.83	15.00
64 GLY O	-16.19	-20.49	57.72	15.00
65 GLY N	-16.79	-20.46	59.86	15.00
65 GLY CA	-17.70	-21.56	59.62	15.00
65 GLY C	-17.39	-22.81	60.42	15.00
65 GLY O	-16.24	-23.13	60.74	15.00
66 GLY N	-18.43	-23.62	60.61	15.00
66 GLY CA	-18.29	-24.85	61.37	15.00
66 GLY C	-19.48	-25.78	61.43	15.00
66 GLY O	-20.57	-25.37	61.00	15.00
67 TYR N	-19.31	-26.94	62.06	15.00
67 TYR CA	-20.37	-27.91	62.16	15.00
67 TYR CB	-19.83	-29.31	61.82	15.00
67 TYR CG	-19.28	-29.38	60.43	15.00
67 TYR CD1	-20.08	-29.14	59.33	15.00
67 TYR CE1	-19.54	-29.09	58.03	15.00
67 TYR CD2	-17.93	-29.57	60.21	15.00
67 TYR CE2	-17.39	-29.52	58.91	15.00
67 TYR CZ	-18.20	-29.27	57.84	15.00
67 TYR OH	-17.70	-29.21	56.59	15.00
67 TYR C	-21.11	-27.86	63.49	15.00
67 TYR O	-20.55	-27.52	64.52	15.00
68 MET N	-22.40	-28.13	63.48	15.00
68 MET CA	-23.12	-28.14	64.76	15.00
68 MET CB	-24.62	-28.23	64.56	15.00
68 MET CG	-25.11	-27.02	63.82	15.00
68 MET SD		-27.31	62.04	15.00
68 MET CE	-26.39		61.56	15.00
68 MET C		-29.31	65.63	15.00
68 MET O	-22.63	-29.21	66.83	15.00
69 THR N	-22.32	-30.42	64.99	15.00
69 THR CA	-21.87	-31.57	65.74	15.00
69 THR CB	-21.55	-32.78	64.81	15.00
69 THR OG1		-32.38	63.72	15.00
69 THR CG2		-33.41	64.29	15.00
69 THR C	-20.65	-31.21	66.61	15.00

69 THR 0	-20.61	-31.62	67.76	15.00
70 ASN N	-19.74	-30.38	66.11	15.00
70 ASN CA	-18.56	-30.05	66.91	15.00
70 ASN CB	-17.53		66.07	15.00
70 ASN CG	-16.83	-30.17	65.02	15.00
70 ASN OD1	-17.07		64.94	15.00
70 ASN ND2	-16.02		64.17	15.00
70 ASN C	-18.99	-29.22	68.11	15.00
70 ASN 0	-18.59	-29.49	69.21	15.00
71 ALA N	-19.95		67.87	15.00
71 ALA CA	-20.42	-27.44	68.91	15.00
71 ALA CB	-21.27	-26.35	68.33	15.00
71 ALA C	-21.15	-28.13	70.05	15.00
71 ALA O	-21.13	-27.69	71.20	15.00
72 PHE N	-21.84	-29.22	69.71	15.00
72 PHE CA	-22.53	-29.99	70.74	15.00
72 PHE CB	-23.59	-30.93	70.13	15.00
72 PHE CG	-24.75	-30.22	69.47	15.00
72 PHE CD1	-25.59	-29.41	70.21	15.00
72 PHE CD2	-25.05	-30.41	68.12	15.00
72 PHE CE1	-26.70	-28.82	69.63	15.00
72 PHE CE2	-26.16	-29.82	67.55	15.00
72 PHE CZ	-26.97	-29.03	68.31	15.00
72 PHE C	-21.49	-30.80	71.51	15.00
72 PHE 0	-21.54	-30.91	72.73	15.00
73 GLN N	-20.55	-31.39	70.78	15.00
73 GLN CA	-19.50	-32.16	71.39	15.00
73 GLN CB	-18.48	-32.63	70.34	15.00
73 GLN CG	-17.59	-33.74	70.84	15.00
73 GLN CD	-17.19	-34.69	69.73	15.00
73 GLN OE1	-17.48	-35.89	69.78	15.00
73 GLN NE2	-16.52	-34.16	68.72	15.00
73 GLN C	-18.81	-31.26	72.43	15.00
73 GLN 0		-31.64	73.59	15.00
74 TYR N	-18.39	-30.06	72.02	15.00
74 TYR CA	-17.70	-29.17	72.97	15.00
74 TYR CB	-17.27	-27.85	72.29	15.00
74 TYR CG	-17.25	-26.64	73.21	15.00
74 TYR CD1	-16.09	-25.24	73.83	15.00
74 TYR CE1	-16.11	-25.21	74.74	15.00
74 TYR CD2		-25.98	73.51	15.00
74 TYR CE2		-24.94	74.43	15.00
4 TYR CZ	-17.30	-24.57	75.05	15.00

74 TYR C -18.51 -28.95 74.27 15 74 TYR O -18.01 -29.19 75.35 15	.00
74 TYR 0 -18.01 -29.19 75.35 15	.00
75 1737 37 10 77	
	.00
75 131 03 00 00 00	.00
75 177 60 00 10	.00
	.00
75 VAL CG1 -23.21 -28.24 75.71 15.	.00
75 VAL CG2 -22.09 -26.67 74.09 15.	00
75 VAL C -20.70 -29.59 76.29 15.	00
75 VAL O -21.29 -29.50 77.38 15.	00
76 GLN N -20.09 -30.71 75.88 15.	00
76 GLN CA -20.03 -31.93 76.68 15.	00
76 GLN CB -20.30 -33.15 75.77 15.	
76 GLN CG -20.03 -34.52 76.35 15.	
76 GLN CD -20.70 -35.58 75.54 15.	
76 GLN OE1 -21.76 -36.09 75.93 15.	
76 GLN NE2 -20.15 -35.88 74.37 15.	
76 GLN C -18.70 -32.10 77.39 15.	
76 GLN O -18.66 -32.13 78.61 15.	
77 LYS N -17.61 -32.20 76.65 15.0	
77 LYS CA -16.31 -32.37 77.28 15.0	
77 LYS CB -15.24 -32.65 76.24 15.0	
77 LYS CG -15.60 -33.84 75.35 15.0	
77 LYS CD -14.38 -34.57 74.81 15.0	
77 LYS CE -13.52 -33.71 73.91 15.0	
77 LYS NZ -12.57 -34.54 73.09 15.0	
77 LYS C -15.94 -31.16 78.15 15.0	
77 LYS 0 -15.31 -31.32 79.19 15.0	
78 ASN N -16.35 -29.96 77.75 15.0	
78 ASN CA -16.09 -28.77 78.57 15.0	
78 ASN CB -16.09 -27.49 77.72 15.0	
78 ASN CG -16.13 -26.22 78.58 15.0	
78 ASN OD1 -15.14 -25.84 79.17 15.0	
78 ASN ND2 -17.32 -25.64 78.72 15.0	
78 ASN C -17.17 -28.68 79.66 15.0	
78 ASN 0 -17.18 -27.77 80.49 15.0	
79 ARG N -18.08 -29.64 79.64 15.0	
79 ARG CA -19.18 -29.69 80.60 15.0	
79 ARG CB -18.69 -30.15 81.98 15.0	
79 ARG CG -18.36 -31.63 82.12 15.00	
79 NRC CD 17 70 00 00	
79 ARG CD -17.79 -31.93 83.52 15.00	
79 ARG CD -17.79 -31.93 83.52 15.00 79 ARG NE -16.63 -32.84 83.47 15.00 79 ARG CZ -15.39 -32.46 83.15 15.00	

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79 ARG NH1			82.86	15.00
79 ARG NH2				
79 ARG C	-20.00			
79 ARG O	-20.27	-27.99		
80 GLY N	-20.32	-27.70		
80 GLY CA	-21.14	-26.50		
80 GLY C	-21.02	-25.34	78.91	
80 GLY O	-19.94		78.31	15.00
81 ILE N	-22.13		78.78	15.00
81 ILE CA	-22.27		77.92	15.00
81 ILE CB	-23.18		76.69	
81 ILE CG2	-24.60		77.13	
81 ILE CG1	-23.28			15.00
81 ILE CD1	-24.04		74.48	15.00
81 ILE C	-22.91		78.75	15.00
81 ILE O	-23.73		79.63	15.00
82 ASP N	-22.58		78.48	15.00
82 ASP CA	-23.13		79.28	15.00
82 ASP CB	-22.13		79.29	15.00
82 ASP CG	-20.88		80.02	15.00
82 ASP OD1	-19.80	-18.87	79.48	15.00
82 ASP OD2	-20.96	-19.62	81.14	15.00
82 ASP C	-24.47	-19.36	78.83	15.00
82 ASP O	-25.10	-19.90	77.94	15.00
83 SER N	-24.92		79.51	15.00
83 SER CA	-26.12	-17.57	79.18	15.00
83 SER CB	-26.70	-16.97	80.48	15.00
83 SER OG	-25.68	-16.38	81.28	15.00
83 SER C	-25.55	-16.45	78.27	15.00
83 SER O	-24.33	-16.24	78.28	15.00
84 GLU N	-26.35	-15.82	77.39	15.00
84 GLU CA	-25.85	-14.72	76.54	15.00
84 GLU CB	-27.00	-13.88	75.94	15.00
84 GLU CG	-27.12	-13.72	74.39	15.00
84 GLU CD	-25.97	-12.97	73.70	15.00
84 GLU OE1	-25.80	-11.74	73.90	15.00
84 GLU OE2	-25.23	-13.64	72.93	15.00
84 GLU C	-25.15	-13.79	77.53	15.00
84 GLU 0		-13.44	77.37	15.00
85 ASP N		-13.43	78.61	15.00
85 ASP CA		-12.53	79.65	15.00
85 ASP CB		-12.68	80.98	15.00
85 ASP CG		-12.06	80.94	
		+2.00	00.34	15.00

85 ASP OD1	_	_	81.40	15.00
85 ASP OD2	-27.63	-10.90		_
85 ASP C	-23.80	-12.67	79.97	
85 ASP O	-23.07		80.00	
86 ALA N	-23.36	-13.90		
86 ALA CA	-21.96	-14.21		
86 ALA CB	-21.89	-15.55		
86 ALA C	-21.00	-14.20	79.40	
86 ALA O	-19.76	-14.22	79.57	
87 TYR N	-21.55		78.20	
87 TYR CA	-20.74	-14.13		15.00
87 TYR CB	-20.41	-15.60	76.66	15.00
87 TYR CG	-18.96	-15.90	76.36	15.00
87 TYR CD1	-18.23	-15.14	75.50	
87 TYR CE1	-16.93	-15.44	75.21	
87 TYR CD2	-18.34	-16.98	76.94	15.00
87 TYR CE2	-17.04		76.65	15.00
87 TYR CZ	-16.35	-16.52	75.79	15.00
87 TYR OH	-15.09	-16.87	75.41	15.00
87 TYR C	-21.57	-13.50	75.86	15.00
87 TYR O	-21.88	-14.19	74.89	15.00
88 PRO N	-21.98	-12.22	75.98	15.00
88 PRO CD	-21.70	-11.28	77.08	15.00
88 PRO CA	-22.78	-11.57	74.93	15.00
88 PRO CB	-22.84	-10.11	75.41	15.00
88 PRO CG	-22.78	-10.21	76.93	15.00
88 PRO C	-22.14	-11.72	73.54	15.00
88 PRO O	-20.91	-11.95	73.44	15.00
89 TYR N	-22.97	-11.62	72.49	15.00
89 TYR CA	-22.50	-11.76	71.11	15.00
89 TYR CB	-23.64	-12.25	70.22	15.00
89 TYR CG	-23.22	-12.67	68.82	15.00
89 TYR CD1	-22.42	-13.82	68.64	15.00
89 TYR CE1	-22.03	-14.24	67.35	15.00
89 TYR CD2	-23.62	-11.92	67.68	15.00
89 TYR CE2	-23.24	-12.31	66.41	15.00
89 TYR CZ	-22.44	-13.47	66.24	15.00
89 TYR OH	-22.00	-13.78	64.96	15.00
89 TYR C	-21.91	-10.45	70.58	15.00
89 TYR O		-9.40	70.70	15.00
90 VAL N	-20.67	-10.51	70.08	15.00
90 VAL CA	-19.98	-9.36	69.50	15.00
90 VAL CB	-18.74	-8.92	70.32	15.00

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90 VAL CG1	-19.17		71.75	15.00
90 VAL CG2	-17.66		70.28	15.00
90 VAL C	-19.58		68.03	15.00
90 VAL O	-18.77	-8.86	67.47	15.00
91 GLY N	-20.12		67.43	15.00
91 GLY CA	-19.85	-10.97	66.03	15.00
91 GLY C	-18.40	-10.89	65.56	15.00
91 GLY O	-18.09		64.36	15.00
92 GLN N	-17.49	-	66.50	15.00
92 GLN CA	-16.08		66.22	15.00
92 GLN CB	-15.44		66.78	15.00
92 GLN CG	-14.07		66.26	15.00
92 GLN CD	-13.74		66.30	15.00
92 GLN OE1	-13.84		65.30	15.00
92 GLN NE2			67.49	15.00
92 GLN C	-15.63		67.00	15.00
92 GLN O	-16.16	-12.45	68.08	15.00
93 GLU N	-14.75	-12.96	66.42	15.00
93 GLU CA	-14.27	-14.11	67.13	15.00
93 GLU CB	-13.67	-15.14	66.19	15.00
93 GLU CG	-13.54	-16.51	66.82	15.00
93 GLU CD	-12.31	-17.23	66.33	15.00
93 GLU OE1	-11.79	-18.11	67.03	15.00
93 GLU OE2	-11.86	-16.90	65.23	15.00
93 GLU C	-13.17	-13.56	68.00	15.00
93 GLU O	-12.58	-12.52	67.67	15.00
94 GLU N	-12.98	-14.19	69.17	15.00
94 GLU CA	-11.95	-13.83	70.12	15.00
94 GLU CB	-12.45	-12.76	71.08	15.00
94 GLU CG	-13.44	-11.78	70.55	15.00
94 GLU CD	-13.36	-10.49	71.28	15.00
94 GLU OE1	-12.35	-9.77	71.16	15.00
94 GLU OE2	-14.28		72.03	15.00
94 GLU C	-11.52	-15.03	70.95	15.00
94 GLU O	-12.21	-1.6.05	71.03	15.00
95 SER N	-10.46	-14.79	71.73	15.00
95 SER CA	-9.89	-15.79	72.60	15.00
95 SER CB	-8.61	-15.24	73.23	15.00
95 SER OG	-7.71	-14.80	72.23	15.00
95 SER C	-10.92	-16.24	73.63	15.00
95 SER O		-15.42	74.18	15.00
96 CYS N		-17.55	73.82	15.00
96 CYS CA	-11.94	-18.19	74.74	15.00

96 CYS C	-11.83	-17.70	76.17	15.00
96 CYS O	-10.75	-17.73	76.79	15.00
96 CYS CB	-11.80	-19.73	74.68	15.00
96 CYS SG	-12.78	-20.81	75.78	15.00
97 MET N	-12.97	-17.26	76.69	15.00
97 MET CA	-13.10	-16.73	78.03	15.00
97 MET CB	-13.17	-15.20	78.02	15.00
97 MET CG	-12.67	-14.49	79.27	15.00
97 MET SD	-10.82	-14.26	79.19	15.00
97 MET CE	-10.31	-15.40	80.50	15.00
97 MET C	-14.34	-17.32	78.74	15.00
97 MET 0	-15.06	-16.60	79.42	15.00
98 TYR N	-14.60	-18.61	78.59	15.00
98 TYR CA	-15.74	-19.28	79.24	15.00
98 TYR CB	-15.63	-20.78	79.00	15.00
98 TYR CG	-16.63	-21.59	79.77	15.00
98 TYR CD1	-17.98	-21.47	79.50	15.00
98 TYR CE1	-18.91	-22.15	80.24	15.00
98 TYR CD2	-16.22	-22.44	80.82	15.00
98 TYR CE2	-17.15	-23.16	81.58	15.00
98 TYR CZ	-18.51	-22.99	81.27	15.00
98 TYR OH	-19.51	-23.63	81.97	15.00
98 TYR C	-15.85	-19.00	80.73	15.00
98 TYR O	-14.85	-18.70	81.38	15.00
99 ASN N	-17.06	-19.08	81.26	15.00
99 ASN CA	-17.25	-18.79	82.67	15.00
99 ASN CB	-17.77	-17.37	82.86	15.00
99 ASN CG	-17.78	-16.95	84.33	15.00
99 ASN OD1	-18.62	-17.40	85.11	15.00
99 ASN ND2	-16.82	-16.11	84.70	15.00
99 ASN C	-18.13	-19.73	83.47	15.00
99 ASN 0	-19.36	-19.70	83.36	15.00
100 PRO N	-17.51	-20.60	84.28	15.00
100 PRO CD	-16.06		84.34	15.00
100 PRO CA		-21.55	85.10	15.00
100 PRO CB		-21.89	86.20	15.00
100 PRO CG	-15.98	-22.00	85.45	15.00
100 PRO C	-19.56	-21.03	85.71	15.00
100 PRO O		21.68		15.00
101 THR N		-19.89		
101 THR CA		-19.30	87.10	15.00
101 THR CB		-18.20	88.13	15.00
101 THR OG1	-19.04	-17.45	87.66	15.00

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		CG2	-19.82	-18.87	89.46	15.00
101		C	-21.77	-18.86	86.17	15.00
101			-22.95	-18.78	86.56	15.00
102	GLY	N	-21.40	-18.68	84.91	15.00
102	GLY	CA	-22.35	-18.33	83.88	15.00
102	GLY	C	-23.27	-19.50	83.58	15.00
102	GLY	. 0	-24.47	-19.30	83.42	15.00
103	LYS	N	-22.68	-20.69	83.38	15.00
103	LYS	CA	-23.39	-21.95	83.13	15.00
103	LYS	CB	-22.95	-23.01	84.16	15.00
103	LYS	CG	-23.73	-24.31	84.12	15.00
103	LYS	CD	-23.11	~25.38	85.05	15.00
103		CE	-21.69	-25.76	84.61	15.00
103		NZ	-21.01	-26.68	85.56	15.00
103			-24.90	-21.82	83.20	15.00
	LYS		-25.45	-21.45	84.23	15.00
	ALA		-25.58	-22.16	82.10	15.00
	ALA		-27.05	-22.11	82.04	15.00
	ALA	_	-27.54	-20.84	81.26	15.00
	ALA		-27.64	-23.39	81.42	15.00
	ALA		-28.85	-23.61	81.42	15.00
	ALA		-26.79	-24.26	80.92	15.00
	ALA		-27.24	-25.50	80.30	15.00
	ALA		-28.05	-25.22	79.03	15.00
105			-26.03	-26.35	79.97	15.00
105			-24.88	-25.90	80.04	15.00
	LYS		-26.32	-27.59	79.60	15.00
	LYS		-25.32	-28.57	79.25	15.00
	LYS		-24.74	-29.13	80.55	15.00
106	LYS		-23.71	-30.22	80.42	15.00
106	LYS		-23.54	-30.96	81.75	15.00
106	LYS		-24.85	-31.60	82.15	15.00
	LYS		-24.61	-32.90	82.82	15.00
	LYS		-26.09	-29.63	78.43	15.00
106		0	-27.32	-29.62	78.34	15.00
107		N	-25.36	-30.51	77.77	15.00
107		CA	-26.00	-31.54	76.97	15.00
107		CB	-26.06	-31.11	75.50	15.00
107		SG	-24.54	-31.49	74.55	15.00
107	CYS	С	-25.07	-32.72	77.10	15.00
107		0	-23.85	-32.54	77.25	15.00
108	ARG		-25.62	-33.92	77.06	15.00
108	ARG	CA	-24.80	-35.12	77.17	15.00

108 ARG CB	-25.28	-36.03	78.28	15.00
108 ARG CG	-25.35	-35.42	79.64	
108 ARG CD	-25.75	-36.49		15.00
108 ARG NE	-27.18	-36.72	80.71	15.00
108 ARG CZ	-27.75	-37.86	81.09	
108 ARG NH1	-27.00			
108 ARG NH2	-29.07		81.22	15.00
108 ARG C	-24.85	-35.89	75.85	15.00
108 ARG O	-25.59		75.72	15.00
109 GLY N	-24.11	-35.40	74.87	15.00
109 GLY CA	-24.05		73.57	15.00
109 GLY C	-24.99		72.55	15.00
109 GLY O	-25.58	-34.38	72.78	15.00
110 TYR N	-25.00	-36.06	71.36	15.00
110 TYR CA	-25.86	-35.67	70.23	15.00
110 TYR CB	-25.19	-34.54		15.00
110 TYR CG	-23.92	-34.95		15.00
110 TYR CD1	-22.72	-34.87	69.39	15.00
110 TYR CE1	-21.60	-35.34	68.80	15.00
110 TYR CD2	-23.95	-35.49	67.52	15.00
110 TYR CE2	-22.82	-35.97	66.92	15.00
110 TYR CZ	-21.64	-35.90	67.55	15.00
110 TYR OH	-20.45	-36.35	66.94	15.00
110 TYR C	-26.23	-36.85	69.29	15.00
110 TYR O	-25.46	-37.79	69.07	15.00
111 ARG N	-27.40	-36.72	68.69	15.00
111 ARG CA	-27.92	-37.70	67.75	15.00
111 ARG CB	-29.27	-38.26	68.26	15.00
111 ARG CG	-29.50	-39.79	68.03	15.00
111 ARG CD	-28.98	-40.22	66.68	15.00
111 ARG NE	-29.52	-41.47	66.17	15.00
111 ARG CZ	-30.81	-41.70	65.92	15.00
111 ARG NH1	-31.74	-40.77	66.15	15.00
111 ARG NH2	-31.18	-42.86	65.41	15.00
111 ARG C	-28.09	-37.07	66.36	15.00
111 ARG O	-28.58	-35.94	66.20	15.00
112 GLU N	-27.59	-37.78	65.36	15.00
112 GLU CA	-27.69	-37.35	63.96	15.00
112 GLU CB	-26.37	-37.55	63.21	15.00
112 GLU CG	-25.23	-36.68	63.74	15.00
112 GLU CD		-36.94	63.06	15.00
112 GLU OE1	-23.53		62.19	15.00
112 GLU OE2	-23.25	-37.90	63.42	15.00

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112 GLU C	-28.84		63.31	15.00
112 GLU 0	-29.00		63.50	15.00
113 ILE N	-29.66		62.57	15.00
113 ILE CA	-30.83		61.85	15.00
113 ILE CB	-31.89		61.73	15.00
113 ILE CG2	-32.92	-37.06	60.63	15.00
113 ILE CG1	-32.57	-36.51	63.08	15.00
113 ILE CD1	-33.32	-35.23	63.13	15.00
113 ILE C	-30.46	-38.42	60.45	15.00
113 ILE 0	-29.74	-37.79	59.70	15.00
114 PRO N	-30.93	-39.62	60.13	15.00
114 PRO CD	-31.77	-40.50	60.95	15.00
114 PRO CA	-30.66	-40.27	58.84	15.00
114 PRO CB	-31.85	-41.21	58.73	15.00
114 PRO CG	-31.86	-41.77	60.07	15.00
114 PRO C	-30.61	-39.34	57.65	15.00
114 PRO O	-31.62	-38.83	57.20	15.00
115 GLU N	-29.44	-39.26	57.05	15.00
115 GLU CA	-29.18	-38.44	55.87	15.00
115 GLU CB	-27.80	-38.78	55.30	15.00
115 GLU CG	-27.56	-38.41	53.84	15.00
115 GLU CD	-26.48	-39.24	53.21	15.00
115 GLU OE1	-26.78	-40.40	52.85	15.00
115 GLU OE2	-25.33	-38.75	53.12	15.00
115 GLU C	-30.20	-38.50	54.78	15.00
115 GLU 0	-30.37	-39.53	54.16	15.00
116 GLY N	-30.90	-37.40	54.58	15.00
116 GLY CA	-31.88	-37.30	53.52	15.00
116 GLY C	-33.29	-37.78	53.79	15.00
116 GLY 0	-34.11	-37.73	52.87	15.00
117 ASN N	-33.56	-38.25	55.01	15.00
117 ASN CA	-34.89	-38.73	55.36	15.00
117 ASN CB	-34.76	-40.00	56.22	15.00
117 ASN CG	-36.06	-40.77	56.33	15.00
117 ASN OD1	-37.13	-40.29	55.89	15.00
117 ASN ND2	-35.99	-41.96	56.91	15.00
117 ASN C	-35.76	-37.69	56.07	15.00
117 ASN 0	-35.55	-37.38	57.23	15.00
118 GLU N	-36.72	-37.13	55.34	15.00
118 GLU CA	-37.65	-36.15	55.89	15.00
118 GLU CB		-35.34	54.78	15.00
118 GLU CG	-37.31	-34.44	54.07	15.00
118 GLU CD	-38.00	-33.61	53.01	15.00

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118 GLU OE1	-38.09	-34.08	51.85	15.00
118 GLU OE2	-38.48	-32.51	53.32	15.00
118 GLU C	-38.73	-36.78	56.75	15.00
118 GLU O	-39.42	-36.09	57.50	15.00
119 LYS N	-38.84	-38.11	56.68	15.00
119 LYS CA	-39.82	-38.82	57.49	15.00
119 LYS CB	-40.11		56.91	15.00
119 LYS CG	-40.84	-41.15	57.86	15.00
119 LYS CD	-41.21	-42.46	57.19	15.00
119 LYS CE	-39.98	-43.29	56.80	15.00
119 LYS NZ	-38.99	-43.60	57.92	15.00
119 LYS C	-39.24	-38.95	58.88	15.00
119 LYS O	-39.97	-38.92	59.86	15.00
120 ALA N	-37.92	-39.06	58.95	15.00
120 ALA CA	-37.26	-39.21	60.24	15.00
120 ALA CB	-35.86	-39.73	60.05	15.00
120 ALA C	-37.20	-37.84	60.87	15.00
120 ALA O	-37.26	-37.70	62.09	15.00
121 LEU N	-37.08	-36.82	60.03	15.00
121 LEU CA	-36.98	-35.45	60.51	15.00
121 LEU CB	-36.44	-34.53	59.41	15.00
121 LEU CG	-36.17	-33.05	59.73	15.00
121 LEU CD1	-35.09	-32.86	60.74	15.00
121 LEU CD2	-35.74	-32.40	58.46	15.00
121 LEU C	-38.30	-34.95	61.07	15.00
121 LEU O	-38.33	-34.36	62.16	15.00
122 LYS N	-39.40	-35.27	60.39	15.00
122 LYS CA	-40.71	-34.88	60.89	15.00
122 LYS CB	-41.80	-35.31	59.90	15.00
122 LYS CG	-43.25	-35.03	60.34	15.00
122 LYS CD	-44.28	-35.77	59.47	15.00
122 LYS CE	-45.67	-35.70	60.08	15.00
122 LYS NZ	-46.60	-36.58	59.32	15.00
122 LYS C	-40.88	-35.54	62.26	15.00
122 LYS 0	-41.33	-34.90	63.22	15.00
123 ARG N	-40.48	-36.81	62.38	15.00
123 ARG CA	-40.58	-37.50	63.65	15.00
123 ARG CB	-40.25	-38.98	63.52	15.00
123 ARG CG	-41.30	-39.84	62.88	15.00
123 ARG CD	-41.30	-41.21	63.51	15.00
123 ARG NE	-39.95	-41.78	63.60	15.00
123 ARG CZ	-39.25	-42.23	62.56	15.00
123 ARG NH1	-39.75	-42.19	61.32	15.00

Section Contraction

123 ARG NH2	~38.05	-42.77	62.77	15.00
123 ARG C	-39.70	-36.88	64.73	15.00
123 ARG O	-40.18	-36.62	65.83	15.00
124 ALA N	-38.45	-36.57	64.41	15.00
124 ALA CA	-37.57	-35.99	65.43	15.00
124 ALA CB	-36.19	-35.74	64.86	15.00
124 ALA C	-38.12	-34.71	66.02	15.00
124 ALA O	-38.00	-34.46	67.22	15.00
125 VAL N	-38.77	-33.92	65.17	15.00
125 VAL CA	-39.36	-32.62	65.52	15.00
125 VAL CB	-39.54	-31.74	64.24	15.00
125 VAL CG1	-40.40	-30.50	64.54	15.00
125 VAL CG2	-38.15	-31.36	63.70	15.00
125 VAL C	-40.70	-32.75	66.24	15.00
125 VAL 0	-41.14	-31.81	66.92	15.00
126 ALA N	-41.40	-33.85	66.02	15.00
126 ALA CA	-42.67	-34.05	66.68	15.00
126 ALA CB	-43.57	-35.00	65.86	15.00
126 ALA C	-42.42	-34.59	68.08	15.00
126 ALA O	-42.99	-34.11	69.06	15.00
127 ARG N	-41.47	-35.52	68.18	15.00
127 ARG CA	-41.15	-36.15	69.47	15.00
127 ARG CB	-40.71	-37.61	69.26	15.00
127 ARG CG	-41.77	-38.56	68.76	15.00
127 ARG CD	-42.88	-38.82	69.77	15.00
127 ARG NE	-43.75	-39.92	69.33	15.00
127 ARG CZ	-44.83	-40.35	69.98	15.00
127 ARG NH1	-45.54	-41.36	69.47	15.00
127 ARG NH2	-45.21	-39.78	71.13	15.00
127 ARG C	-40.13	-35.44	70.36	15.00
127 ARG 0	-40.28	-35.42	71.58	15.00
128 VAL N	-39.07	-34.89	69.76	15.00
128 VAL CA	-37.97	-34.24	70.49	15.00
128 VAL CB	-36.63	-34.58	69.81	15.00
128 VAL CG1	-35.45	-34.03	70.61	15.00
128 VAL CG2	-36.48	-36.07	69.66	15.00
128 VAL C	-38.10	-32.73	70.56	15.00
128 VAL 0	-37.92	-32.12	71.63	15.00
129 GLY N	-38.44	-32.14	69.42	15.00
129 GLY CA	-38.57	-30.70	69.33	15.00
129 GLY C	-37.60	-30.09	68.33	15.00
129 GLY O	-37.12	-30.79	67.45	15.00
130 PRO N	-37.31	-28.77	68.42	15.00

130 PRO CD	-37.83	-27.89	69.47	15.00
130 PRO CA	-36.40	-28.02	67.54	15.00
130 PRO CB	-36.13	-26.77	68.34	15.00
130 PRO CG	-37.40	-26.52	68.98	15.00
130 PRO C	-35.13	-28.78	67.19	15.00
130 PRO O	-34.38	-29.24	68.05	15.00
131 VAL N	-34.91	-28.91	65.89	15.00
131 VAL CA	-33.78	-29.63	65.37	15.00
131 VAL CB	-34.27	-30.83	64.56	15.00
131 VAL CG1	-33.10	-31.52	63.87	15.00
131 VAL CG2	-35.02	-31.76	65.47	15.00
131 VAL C	-32.91	-28.75	64.48	15.00
131 VAL 0	-33.43	-28.02	63.65	15.00
132 SER N	-31.60	-28.85	64.68	15.00
132 SER CA	-30.59	-28.13	63.91	15.00
132 SER CB	-29.21	-28.32	64.54	15.00
132 SER OG	-28.89	-27.37	65.52	15.00
132 SER C	-30.50	-28.73	62.50	15.00
132 SER O	-30.34	-29.94	62.34	15.00
133 VAL N	-30.47	-27.87	61.50	15.00
133 VAL CA	-30.38	-28.36	60.13	15.00
133 VAL CB	-31.75	-28.35	59.45	15.00
133 VAL CG1	-32.73	-29.24	60.17	15.00
133 VAL CG2	-32.30	-26.98	59.41	15.00
133 VAL C	-29.52	-27.41	59.35	15.00
133 VAL O	-29.29	-26.29	59.79	15.00
134 ALA N	-29.14	-27.85	58.15	15.00
134 ALA CA	-28.32	-27.08	57.22	15.00
134 ALA CB	-26.99	-27.75	56.97	15.00
134 ALA C	-29.14	-27.06	55.95	15.00
134 ALA O	-29.99	-27.92	55.71	15.00
135 ILE N	-28.79	-26.11	55.09	15.00
135 ILE CA	-29.49	-25.93	53.83	15.00
135 ILE CB		-25.12	54.04	15.00
135 ILE CG2		-26.00		
135 ILE CG1		-23.83	54.85	15.00
135 ILE CD1	-31.76	-22.91	54.86	15.00
135 ILE C		-25.16	52.82	15.00
135 ILE O	-27.58	-24.73	53.10	15.00
136 ASP N		-25.10	51.61	15.00
136 ASP CA	-28.56			
136 ASP CB	-28.74			
136 ASP CG	-28.23	-24.06	48.08	15.00

136 ASP OD1	-28.28	-24.45	46.91	15.00
136 ASP OD2	-27.73	-22.96		15.00
136 ASP C	-29.32	-23.01	50.62	15.00
136 ASP O	-30.51		50.35	15.00
137 ALA N	-28.59	-21.94	50.84	15.00
137 ALA CA	-29.23		50.97	15.00
137 ALA CB	-29.22		52.43	15.00
137 ALA C	-28.65		50.07	15.00
137 ALA O	-28.89	-18.38	50.30	15.00
138 SER N	-27.97	-19.97	49.00	15.00
138 SER CA	-27.34			15.00
138 SER CB	-26.28		47.23	15.00
138 SER OG	-26.71	-21.14	47.02	15.00
138 SER C	-28.32	-18.41	47.07	15.00
138 SER O	-28.09		46.57	15.00
139 LEU N	-29.42		46.81	15.00
139 LEU CA	-30.44		45.87	15.00
139 LEU CB	-31.60		45.83	15.00
139 LEU CG	-31.57	-20.77	44.76	15.00
139 LEU CD1	-31.68	-20.14	43.39	15.00
139 LEU CD2	-30.29		44.84	15.00
139 LEU C	-30.95		46.25	15.00
139 LEU O	-31.39	-17.09	47.37	15.00
140 THR N	-30.99	-16.39	45.28	15.00
140 THR CA	-31.41	-15.04	45.54	15.00
140 THR CB	-31.16	-14.10	44.30	15.00
140 THR OG1	-30.83	-14.87	43.13	15.00
140 THR CG2	-30.00	-13.22	44.59	15.00
140 THR C	-32.86	-14.97	46.00	15.00
140 THR O	-33.25	-14.00	46.66	15.00
141 SER N	-33.65	-15.99	45.68	15.00
141 SER CA	-35.05	-16.03	46.09	15.00
141 SER CB	-35.80	-17.14	45.35	15.00
141 SER OG	-34.95	-18.27	45.17	
141 SER C	-35.15	-16.16	47.60	
141 SER O	-35.95	-15.48	48.25	15.00
142 PHE N	-34.23	-16.95	48.15	15.00
142 PHE CA	-34.11	-17.19	49.58	15.00
142 PHE CB	-32.92	-18.13	49.84	15.00
142 PHE CG	-32.94	-18.73		15.00
142 PHE CD1		-20.03		15.00
142 PHE CD2		-17.97		15.00
142 PHE CE1		-20.55		15.00
				_

142 PHE CE2	-32 62	10.40	53.60	
142 PHE CZ	-33.10	-18.49		15.00
142 PHE C	-33.10		53.78	15.00
142 PHE 0	-34.57		50.34	15.00
143 GLN N	-33.02	-15.60	51.33	15.00
143 GLN CA	-32.74	-15.01	49.84	15.00
143 GLN CB	-31.45			15.00
143 GLN CG	-30.34		49.95	15.00
143 GLN CD	-29.07		49.75	15.00
143 GLN OE1	-29.07		49.24	15.00
143 GLN NE2	-29.02		49.08	15.00
143 GLN C	-33.90		49.00	15.00
143 GLN 0	-34.20		50.35	15.00
143 GHN 0	-34.58	-12.02	51.29	15.00
144 PHE CA	-35.72	-12.77	49.20	15.00
144 PHE CB	-35.72	-11.87 -11.31	49.00	15.00
144 PHE CG	-34.56		47.56	15.00
144 PHE CD1	-34.12		47.21	15.00
144 PHE CD2	-33.89		48.08	15.00
144 PHE CE1	-33.03	-8.71	46.02	15.00
144 PHE CE2	-32.80		47.77	15.00
144 PHE CZ	-32.36	-8.93	45.68 46.55	15.00
144 PHE C	-37.07	-12.48	49.36	15.00
144 PHE 0	-38.09	-11.81	49.24	15.00 15.00
145 TYR N	-37.08	-13.74	49.80	15.00
145 TYR CA	-38.31	-14.45	50.22	15.00
145 TYR CB	-37.94	-15.77	50.96	15.00
145 TYR CG	-39.07	-16.36	51.80	15.00
145 TYR CD1	-39.94	-17.34	51.29	15.00
145 TYR CE1	-41.00	-17.82	52.05	15.00
145 TYR CD2	-39.30		53.10	15.00
145 TYR CE2	-40.33	-16.32	53.86	15.00
145 TYR CZ	-41.19	-17.29	53.34	15.00
145 TYR OH	-42.25	-17.67		15.00
145 TYR C		-13.55		15.00
145 TYR 0	-38.60	-12.81	51.97	15.00
146 SER N	-40.45	-13.64	51.02	15.00
146 SER CA	-41.31	-12.83	51.88	15.00
146 SER CB	-41.74	-11.55	51.16	15.00
146 SER OG	-40.94	-10.40	51.51	15.00
146 SER C	-42.53	-13.55	52.45	15.00
146 SER O	-42.89	-13.33	53.61	15.00
147 LYS N	-43.15	-14.44	51.68	15.00

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147 LYS CA	-44.33	-15.12	52.17	15.00
147 LYS CB	-45.53	-14.22		15.00
147 LYS CG	-45.80	-13.23	53.02	15.00
147 LYS CD	-46.76	-12.13		15.00
147 LYS CE	-47.91	-12.66		15.00
147 LYS NZ	-48.65	-13.77	52.35	15.00
147 LYS C	-44.63	-16.41	51.41	15.00
147 LYS O	-44.45	-16.45	50.20	15.00
148 GLY N	-45.09	-17.43	52.10	15.00
148 GLY CA	-45.45	-18.66	51.42	15.00
148 GLY C	-44.53	-19.82	51.65	15.00
148 GLY O	-43.56	-19.73	52.41	15.00
149 VAL N	-44.79	-20.91	50.92	15.00
149 VAL CA	-43.99	-22.13	50.98	15.00
149 VAL CB	-44.85	-23.39	50.69	15.00
149 VAL CG1	-43.98	-24.66	50.78	15.00
149 VAL CG2	-45.98	-23.43	51.62	15.00
149 VAL C	-42.91	-22.02	49.90	15.00
149 VAL O	-43.22	-22.06	48.72	15.00
150 TYR N	-41.68	-21.79	50.34	15.00
150 TYR CA	-40.49	-21.66	49.49	15.00
150 TYR CB	-39.31	-21.17	50.33	15.00
150 TYR CG	-38.07	-21.02	49.48	15.00
150 TYR CD1	-37.90	-19.91	48.65	15.00
150 TYR CE1	-36.75	-19.75	47.90	15.00
150 TYR CD2	-37.05	-21.97	49.52	15.00
150 TYR CE2	-35.89	-21.82	48.77	15.00
150 TYR CZ	-35.75	-20.71	47.97	15.00
150 TYR OH	-34.59	-20.55	47.26	15.00
150 TYR C	-40.04	-22.89	48.73	15.00
150 TYR O	-39.72	-23.95	49.29	15.00
151 TYR N	-39.88	-22.68	47.44	15.00
151 TYR CA	-39.46	-23.73	46.54	15.00
151 TYR CB	-40.63	-24.64	46.18	15.00
151 TYR CG	-40.18	-25.91	45.48	15.00
151 TYR CD1		-26.99	46.24	15.00
151 TYR CE1		-28.12	45.59	15.00
151 TYR CD2		-25.99	44.05	15.00
151 TYR CE2		-27.07	43.37	15.00
151 TYR CZ		-28.15	44.13	15.00
151 TYR OH		-29.25	43.41	15.00
151 TYR C		-23.0 <b>9</b>	45.29	15.00
151 TYR O	-39.45	-22.21	44.68	15.00

152 ASP N	-37.66	-23.52	44.96	15 00
152 ASP CA	-36.90			
152 ASP CB	-35.90			
152 ASP CG	-35.26		43.08	15.00
152 ASP OD1	-35.42	-20.01	42.99	15.00
152 ASP OD2	-34.63	-21.92		15.00
152 ASP C	-36.16			
152 ASP 0	-35.56		·	15.00
153 GLU N	-36.16			15.00
153 GLU CA	-35.56		41.38	15.00
153 GLU CB	-36.09			15.00 15.00
153 GLU CG	-35.94			15.00
153 GLU CD	-36.44		37.83	15.00
153 GLU OE1	-35.63		36.94	15.00
153 GLU OE2	-37.62	-24.87	37.62	15.00
153 GLU C	-34.04		41.31	
153 GLU O	-33.47		40.99	15.00
154 SER N	-33.39	-24.59	41.60	15.00
154 SER CA	-31.95	-24.48	41.56	15.00
154 SER CB	-31.59	-23.07	41.09	15.00
154 SER OG	-32.22	-22.78	39.85	15.00
154 SER C	-31.28	-24.79	42.91	15.00
154 SER O	-30.07	-24.67	43.03	15.00
155 CYS N	-32.08	-25.19	43.90	15.00
155 CYS CA	-31.55	-25.49	45.23	15.00
155 CYS C	-30.88	-26.84	45.21	15.00
155 CYS O	<b>-</b> 31.52	-27.87	45.08	15.00
155 CYS CB	-32.63	-25.43	46.33	15.00
155 CYS SG	-32.25	-24.44	47.82	15.00
156 ASN N	-29.57	-26.79	45.39	15.00
156 ASN CA	-28.70	-27.97	45.38	15.00
156 ASN CB	-27.30			15.00
156 ASN CG 156 ASN OD1	-26.51	-28.59	44.35	15.00
	-26.70		44.62	15.00
156 ASN ND2 156 ASN C	-25.58		43.51	15.00
156 ASN C			46.72	15.00
157 SER N		-28.19	47.70	15.00
157 SER N		-29.96	46.73	15.00
157 SER CA		~30.77	47.95	15.00
157 SER CB		-32.11	47.70	
157 SER OG 157 SER C		-31.95	47.10	
157 SER C	-27.68 -27.48			15.00
TO! SEK O	27.48	-31.52	49.54	15.00

158 ASP N	-26.71	-30.93	47.55	15.00
158 ASP CA	-25.34	-31.23		
158 ASP CB	-24.69	-32.07		
158 ASP CG	-25.32	-33.44	46.67	
158 ASP OD1	-26.15	-33.63		
158 ASP OD2	-25.05			
158 ASP C	-24.46	-30.07		
158 ASP 0	-23.27	-30.25		15.00
159 ASN N	-25.05	-28.88		
159 ASN CA	-24.30			15.00
159 ASN CB	-24.34			15.00
159 ASN CG	-23.51		47.62	15.00
159 ASN OD1	-22.35		48.00	15.00
159 ASN ND2	-24.12	-24.33	47.38	15.00
159 ASN C	-24.93		49.82	
159 ASN 0	-25.67	-26.05		15.00
160 LEU N	-24.68	-27.59	50.99	15.00
160 LEU CA	-25.16	-27.07	52.24	15.00
160 LEU CB	-25.06	-28.15	53.29	15.00
160 LEU CG	-25.72		52.83	15.00
160 LEU CD1	-25.24		53.62	15.00
160 LEU CD2	-27.22	-29.26	52.91	15.00
160 LEU C	-24.22	-25.95	52.63	15.00
160 LEU O	-23.06	-26.21	52.92	15.00
161 ASN N	-24.73	-24.74	52.74	15.00
161 ASN CA	-23.91	-23.58	53.10	15.00
161 ASN CB	-23.61	-22.78	51.83	15.00
161 ASN CG	-24.84	-22.55	50.93	15.00
161 ASN OD1	-25.71	-21.74	51.22	15.00
161 ASN ND2	-24.92	-23.31	49.84	15.00
161 ASN C	-24.42	-22.67	54.23	15.00
161 ASN 0	-23.75	-21.70	54.59	15.00
162 HIS N		-22.96	54.76	15.00
162 HIS CA	-26.13	-22.19	55.86	15.00
162 HIS CB		-21.15	55.35	15.00
162 HIS CG	-27.63	-20.21	56.41	15.00
162 HIS CD2	-28.89	-19.99	56.84	15.00
162 HIS ND1	-26.82	-19.39	57.16	15.00
162 HIS CE1	-27.56	-18.68	57.99	15.00
162 HIS NE2	-28.82	-19.03	57.82	15.00
162 HIS C	-26.80	-23.11	56.86	15.00
162 HIS O	-27.41	-24.10	56.49	15.00
163 ALA N	-26.61	-22.84	58.14	15.00

169 TYR CZ	-45.12		62.55	15.00
169 TYR OH	-45.63	-33.71	63.41	15.00
169 TYR C	-45.53	-28.26	60.32	15.00
169 TYR 0	-46.20	-28.25	61.37	15.00
170 GLY N	-46.09	-28.23	59.12	15.00
170 GLY CA	-47.54	-28.17	58.95	15.00
170 GLY C	-47.93	-28.06	57.48	15.00
170 GLY 0	-47.12	-28.35	56.61	15.00
171 ILE N	-49.14	-27.61	57.17	15.00
171 ILE CA	-49.55	-27.54	55.77	15.00
171 ILE CB	-50.30	-28.85	55.41	15.00
171 ILE CG2	-51.37	-29.17	56.40	15.00
171 ILE CG1	-50.79	-28.82	53.99	15.00
171 ILE CD1	-51.11	-30.20	53.45	15.00
171 ILE C	-50.38	-26.29	55.43	15.00
171 ILE 0	-51.37	-26.02	56.10	15.00
172 GLN N	-49.93	-25.52	54.44	15.00
172 GLN CA	-50.64	-24.31	54.01	15.00
172 GLN CB	-49.68	-23.10	54.05	15.00
172 GLN CG	-50.30	-21.69	53.87	15.00
172 GLN CD	-49.27	-20.56	53.75	15.00
172 GLN OE1	-48.07	-20.77	53.89	15.00
172 GLN NE2	~49.74	-19.35	53.49	15.00
172 GLN C	-51.25	-24.41	52.63	15.00
172 GLN O	-50.56	-24.19	51.64	15.00
173 LYS N	-52.55	-24.75	52.57	15.00
173 LYS CA	~53.33	-24.85	51.32	15.00
173 LYS CB	-53.40	-23.48	50.61	15.00
173 LYS CG	-54.16	-22.39	51.36	15.00
173 LYS CD	-53.57	-22.02	52.75	15.00
173 LYS CE	-54.37	-22.65	53.92	15.00
173 LYS NZ	-54.04	-22.06	55.23	15.00
173 LYS C	-52.85	-25.93	50.36	15.00
173 LYS O	-52.73	-25.69	49.15	15.00
174 GLY N	-52.61	-27.11	50.90	15.00
174 GLY CA	-52.15	-28.21	50.08	15.00
174 GLY C	-50.64	-28.26	50.03	15.00
174 GLY O	-50.08	-29.23	49.53	15.00
175 ASN N		-27.25	50.58	15.00
175 ASN CA		-27.27	50.57	15.00
175 ASN CB		-26.00	49.95	15.00
175 ASN CG		-25.73	48.57	15.00
175 ASN OD1	-48.53	-26.61	47.73	15.00
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61.63

60.20

15.00

15.00

**-44.37 -24.29** 

**-39.62 -23.75** 

179 ILE CD1

179 ILE C

58.11

57.20

54.84

54.04

15.00

15.00

15.00

15.00

-28.66 -14.61

-28.06 -15.46

-28.59 -16.14

-30.63 -15.04

184 TRP CD1

184 TRP NE1

184 TRP CZ2

184 TRP CZ3

WO 97/16177		TABLE V	m	
184 TRP CH2	-29.52	-15.92	53.84	15.00
184 TRP C	-31.34	-11.59	60.02	15.00
184 TRP O	-31.41	-10.70	59.18	15.00
185 GLY N	-31.95	-11.51	61.20	15.00
185 GLY CA	-32.75	-10.34	61.52	15.00
185 GLY C	-34.24	-10.61	61.41	15.00
185 GLY O	-34.63	-11.61	60.82	15.00
186 GLU N	-35.09	-9.75	61.96	15.00
186 GLU CA	-36.52	-10.00	61.83	15.00
186 GLU CB	-37.32	-9.44	63.01	15.00
186 GLU CG	-36.65	-9.45	64.34	15.00
186 GLU CD	-37.34	-8.56	65.32	15.00
186 GLU OE1	-36.68	-8.08	66.25	15.00
186 GLU OE2	-38.55		65.21	15.00
186 GLU C	~36.99	-9.30	60.56	15.00
186 GLU O	-38.10	-9.52	60.08	15.00
187 ASN N	-36.14	-8.44	60.02	15.00
187 ASN CA	-36.46	-7.68	58.83	15.00
187 ASN CB	-35.60	-6.42		15.00
187 ASN CG	-35.84		59. <b>9</b> 7	15.00
187 ASN OD1	-34.92		60.44	15.00
187 ASN ND2	-37.10		60.45	15.00
187 ASN C	-36.21	-8.52	57.60	15.00
187 ASN 0	-36.21	-8.00	56.49	15.00
188 TRP N	-35.89		57.81	15.00
188 TRP CA	-35.66	-10.68	56.68	15.00
188 TRP CB	-34.39	-11.49	56.84	15.00
188 TRP CG	-34.20	-12.53	55.78	15.00
188 TRP CD2	-34.69			15.00
188 TRP CE2	-34.21			15.00
188 TRP CE3 188 TRP CD1		-14.67	56.67	15.00
188 TRP NE1	-33.48	-12.37	54.63	15.00
188 TRP CZ2		-13.54	53.91	15.00
188 TRP CZ3		-15.96		
188 TRP CH2		-16.55	56.35	15.00
188 TRP C		-11.63	55.16 56.71	15.00
188 TRP 0		-11.63	56.71	15.00
189 GLY N		-12.09		15.00
189 GLY CA		-13.05	55.42	15.00
189 GLY C		-13.05	56.18	15.00 15.00
189 GLY O		-11.50		
100 301 0	40.00		56.32	15.00

-40.23 -13.64

190 ASN N

56.73

**-39.11 -17.29** 

-39.23 -16.35

**-39.08 -18.56** 

**-39.32 -18.92** 

-37.98 -18.83

**-36.90 -19.78** 

-38.23 -19.04

**-36.98 -19.00** 

**-39.99 -20.30** 

**-39.67 -21.15** 

-41.01 -20.45

**-41.71 -21.72** 

-43.20 -21.48

58.14

57.35

57.75

56.37

55.52

56.01

54.02

53.22

56.33

57.18

55.48

55.30

55.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

193 TYR C

193 TYR O

194 ILE N

194 ILE CA

194 ILE CB

194 ILE CG2

194 ILE CG1

194 ILE CD1

194 ILE C

194 ILE 0

195 LEU N

195 LEU CA

195 LEU CB

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195 LEU CG	-43.96	-20.94	56.22	15.00
195 LEU CD1	-45.35			
195 LEU CD2	-44.03	-22.03	57.28	15.00
195 LEU C	-41.08	-22.46	54.14	15.00
195 LEU O	-41.24	-22.03	52.99	15.00
196 MET N	-40.38	-23.55		15.00
196 MET CA	-39.73	-24.28	53.34	15.00
196 MET CB	-38.30	-24.61	53.74	15.00
196 MET CG	-37.40	-23.37	53.79	15.00
196 MET SD	-35.72	-23.74	54.33	15.00
196 MET CE	-35.19	-24.86	53.02	15.00
196 MET C	-40.45	-25.54	52.91	15.00
196 MET O	-40.92	-26.29	53.76	15.00
197 ALA N	-40.52	-25.77	51.61	15.00
197 ALA CA	-41.19	-26.95	51.06	15.00
197 ALA CB	-40.98	-26.99	49.58	15.00
197 ALA C	-40.77	-28.28	51.69	15.00
197 ALA O	-39.59	-28.51	51.93	15.00
198 ARG N	-41.77	-29.13	51.96	15.00
198 ARG CA	-41.53		52.55	15.00
198 ARG CB	-42.18	-30.53	53.94	15.00
198 ARG CG	-42.26		54.56	15.00
198 ARG CD	-42.45		56.05	15.00
198 ARG NE	-43.59		56.41	15.00
198 ARG CZ	-44.85	-31.46	56.47	15.00
198 ARG NH1	-45.11	-32.72	56.19	15.00
198 ARG NH2	-45.84		56.84	15.00
198 ARG C	-42.05	-31.55	51.67	15.00
198 ARG O	-43.07	-31.41	50.99	15.00
199 ASN N		-32.69	51.77	15.00
199 ASN CA	-41.70			
199 ASN CB	-43.17		51.14	15.00
199 ASN CG	-43.54	-34.74	52.54	15.00
199 ASN OD1	-42.66	-34.97	53.39	15.00
199 ASN ND2	-44.84	-34.86	52.79	15.00
199 ASN C	-41.32	-33.73	49.56	15.00
199 ASN O	-41.24	-34.72	48.84	15.00
200 LYS N	-41.04	-32.51	49.14	15.00
200 LYS CA		-32.31	47.77	15.00
200 LYS CB		-30.88	47.33	15.00
200 LYS CG	-42.41	-30.50	47.40	15.00
200 LYS CD	-42.69	-29.14	46.77	15.00
200 LYS CE	-42.63	-29.16	45.28	15.00

-30.85

56.66

15.00

-32.59

206 ILE 0

					_	
207	ALA	N	-32.81	-32.22	54.92	15.00
207	ALA	CA	-32.67	-33.38	55.79	15.00
207	ALA	CB	-33.80	-34.33	55.58	15.00
207	ALA	С	-31.34	-34.09	55.63	15.00
207	ALA	0	-31.17	-35.20	56.13	15.00
208	ASN	N	-30.36	-33.45	55.00	15.00
208	ASN	CA	-29.04	-34.07	54.76	15.00
208	ASN	CB	-28.34	-33.45	53.52	15.00
208	ASN	CG	-28.78	-34.07	52.18	15.00
208	ASN	OD1	-29.25	-35.21	52.11	15.00
208	ASN	ND2	-28.60	-33.31	51.12	15.00
208	ASN	С	-28.08	-34.01	55.94	15.00
208	ASN	0	-27.28	-34.93	56.15	15.00
209	LEU	N	-28.08	-32.88	56.64	15.00
209	LEU	CA	-27.19	-32.64	57.78	15.00
209	LEU	CB	-26.07	-31.64	57.40	15.00
209	LEU	CG	-24.73	-31.69	58.15	15.00
209	LEU	CD1	-24.11	-33.07	58.09	15.00
209	LEU	CD2	-23.76	-30.66	57.52	15.00
209	LEU	С	-27.97	-32.12	59.02	15.00
209	LEU	0	-27.72	-30.99	59.50	15.00
210	ALA	N	-28.79	-32.99	59.65	15.00
210	ALA	H	-29.19	-33.47	58.90	15.00
210	ALA	CA	-29.59	-32.51	60.77	15.00
210	ALA	CB	-31.08	-32.79	60.54	15.00
210	ALA	С	-29.19	-33.24	62.06	15.00
210	ALA	0	-28.91	-34.44	62.09	15.00
211	SER	N	-29.24	-32.48	63.17	15.00
211	SER	CA	-28.94	-33.06	64.47	15.00
211	SER	CB	-27.44	-33.22	64.70	15.00
211	SER	OG	-26.78	-31.96	64.75	15.00
211	SER		-29.57	-32.31	65.62	15.00
	SER		-30.17	-31.24	65.41	15.00
	PHE	N	-29.43	-32.92	66.81	15.00
212			-29.96	-32.40	68.07	15.00
212	PHE		-31.46	-32.75	68.27	15.00
212	PHE	CG	-31.78	-34.22	68.27	15.00
212		CD1	-32.33	-34.84	67.14	15.00
212	PHE	CD2	-31.61	-34.97	69.41	15.00
212	PHE			-36.19	67.17	15.00
212	PHE	CE2		-36.30	69.45	15.00
212	PHE	CZ	-32.54	-36.91	68.33	15.00
212	PHE	С	-29.16	-32.88	69.27	15.00

44.32

15.00

-44.49 -25.37

232 HOH OH2

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46.91

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (A2) for the cathepsin K complex with inhibitor 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone.

Residue Atom	x	Y	z	В
1 ALA CB	-54.23	-33.20	65.64	15.00
1 ALA C	-53.54	-33.00	63.24	15.00
1 ALA O	-52.73	-33.83	62.79	15.00
1 ALA N	-54.89	-34.90	63.97	15.00
1 ALA CA	-54.65	-33.45	64.19	15.00
2 PRO N	-53.49	-31.70	62.91	15.00
2 PRO CD	-54.26	-30.56	63.46	15.00
2 PRO CA	-52.44	-31.22	62.00	15.00
2 PRO CB	-52.58	-29.69	62.07	15.00
2 PRO CG	-53.25	-29.45	63.41	15.00
2 PRO C	-51.07	-31.68	62.50	15.00
2 PRO O	-50.74	-31.50	63.67	15.00
3 ASP N	-50.31	-32.35	61.64	15.00
3 ASP CA	-48.98	-32.84	62.01	15.00
3 ASP CB	-48.55	-33.98	61.08	15.00
3 ASP CG	-47.66	-35.01	61.77	15.00
3 ASP OD1	-46.52	-34.66	62.18	15.00
3 ASP OD2	-48.10	-36.17	61.89	15.00
3 ASP C	-48.00	-31.68	61.94	15.00
3 ASP O	-47.13	-31.65	61.06	15.00
4 SER N	-48.12	-30.72	62.85	15.00
4 SER CA	-47.23	-29.56	62.86	15.00
4 SER CB	-47.87	-28.37	62.15	15.00
4 SER OG	-49.05	-27.95	62.82	15.00
4 SER C	-46.76	-29.14	64.25	15.00
4 SER O	-47.54	-29.19	65.22	15.00
5 VAL N	-45.50	-28.70	64.35	15.00
5 VAL CA	-44.91	-28.26	65.61	15.00
5 VAL CB	-43.77	-29.21	66.10	15.00
5 VAL CG1	-43.35	-28.82	67.51	15.00
5 VAL CG2	-44.19	-30.65	66.04	15.00
5 VAL C	-44.29	-26.87	65.45	15.00
5 VAL O	-43.47	-26.65	64.55	15.00
6 ASP N	-44.64	-25.95	66.33	15.00

6 ASP CA	-44.10		66.28	15.00
6 ASP CB	-45.17	-23.57	65.90	15.00
6 ASP CG	-44.59	-22.26	65.40	15.00
6 ASP OD1	-43.41	-21.95	65.68	15.00
6 ASP OD2	-45.34	-21.52	64.73	15.00
6 ASP C	-43.52	-24.32	67.66	15.00
6 ASP O	-44.20	-23.78	68.53	15.00
7 TYR N	-42.27		67.85	15.00
7 TYR CA	-41.62	-24.50	69.13	15.00
7 TYR CB	-40.20	-25.08	69.12	15.00
7 TYR CG	-40.20	-26.58	69.20	15.00
7 TYR CD1	-40.68	-27.24	70.33	15.00
7 TYR CE1	-40.76	-28.62	70.38	15.00
7 TYR CD2	-39.81	-27.36	68.11	15.00
7 TYR CE2	-39.89	-28.74	68.15	15.00
7 TYR CZ	-40.37	-29.36	69.29	15.00
7 TYR OH	-40.51	-30.72	69.31	15.00
7 TYR C	-41.63	-23.07	69.62	15.00
7 TYR O	-41.57	-22.84	70.83	15.00
8 ARG N	-41.74	-22.11	68.70	15.00
8 ARG CA	-41.77	-20.68	69.08	15.00
8 ARG CB	-41.86	-19. <b>7</b> 7	67.84	15.00
8 ARG CG	-40.77	-19.98	66.80	15.00
8 ARG CD	-41.01	-19.12	65. <b>58</b>	15.00
8 ARG NE	-42.34	-19.33	65.02	15.00
8 ARG CZ	-42.70	-18.96	63.80	15.00
8 ARG NH1	-41.83	-18.36	63.00	15.00
8 ARG NH2	-43.94	-19.18	63.38	15.00
8 ARG C	-42.94	-20.39	70.02	15.00
8 ARG O	-42.79	-19.67	71.02	15.00
9 LYS N	-44.10	-20.98	69.72	15.00
9 LYS CA	-45.29	-20.80	70.53	15.00
9 LYS CB	-46.53	-21.26	69.78	15.00
9 LYS CG	-46.84	-20.43	68.56	15.00
9 LYS CD	-48.15	-20.86	67.92	15.00
9 LYS CE	-48.39	-20.11	66.62	15.00
9 LYS NZ	-49.58	-20.62	65.88	15.00
9 LYS C	-45.18	-21.53	71.85	15.00
9 LYS O	-45.95	-21.29	72.77	15.00
10 LYS N	-44.18	-22.40	71.95	15.00
10 LYS CA	-43.99	-23.17	73.16	15.00
10 LYS CB	-43.71	-24.63	72.80	15.00

**-35.29 -13.83** 

-35.03 -12.54

-35.58 -12.84

-34.27 -14.02

15 PRO CA

15 PRO CB

15 PRO CG

15 PRO C

15.00

15.00

15.00

15.00

15.00

70.27

71.03

72.37

15 PRO O	-33.20	-14.57	69.36	15.00
16 VAL N	-34.63	-13.62	67.92	15.00
16 VAL CA	-33.72	-13.74	66.79	15.00
16 VAL CB	-34.41	-13.41	65.45	15.00
16 VAL CG1	-33.40	-13.43	64.31	15.00
16 VAL CG2	-35.52	-14.40	65.18	15.00
16 VAL C	-32.53	-12.82	67.01	15.00
16 VAL O	-32.69	-11.69	67.46	15.00
17 LYS N	-31.34	-13.30	66.69	15.00
17 LYS CA	-30.16	-12.50	66.88	15.00
17 LYS CB	-29.27	-13.11	67.95	15.00
17 LYS CG	-29. <b>9</b> 7	-13.18	69.30	15.00
17 LYS CD	-29.17	-13.95	70.33	15.00
17 LYS CE	-29.96	-14.12	71.61	15.00
17 LYS NZ	-31.23	-14.87	71.36	15.00
17 LYS C	-29.41	-12.30	65.58	15.00
17 LYS 0	-29.51	-13.11	64.66	15.00
18 ASN N	-28.68	-11.18	65.51	15.00
18 ASN CA	-27.89	-10.84	64.33	15.00
18 ASN CB	-28.01	-9.37	63.99	15.00
18 ASN CG	-27.30	-9.03	62.72	15.00
18 ASN OD1	-27.10	-9.90	61.88	15.00
18 ASN ND2	-26.89	-7.78	62.57	15.00
18 ASN C	-26.42	-11.21	64.52	15.00
18 ASN O	-25.77	-10.70	65.43	15.00
19 GLN N	-25.89	-12.04	63.63	15.00
19 GLN CA	-24.50	-12.46	63.73	15.00
19 GLN CB	-24.24	-13.75	62.96	15.00
19 GLN CG	-24.32	-13.63	61.47	15.00
19 GLN CD	-24.07	-14.96	60.80	15.00
19 GLN OE1	-25.01	-15.66	60.43	15.00
19 GLN NE2	-22.81	-15.32	60.65	15.00
19 GLN C	-23.48	-11.40	63.36	
19 GLN O	-22.32	-11.48	63.76	15.00
20 GLY N	-23.90	-10.40	62.59	15.00
20 GLY CA	-22.99	-9.33	62.22	15.00
20 GLY C	-21.93	-9.72	61.22	15.00
20 GLY 0	-22.06	-10.71	60.52	15.00
21 GLN N	-20.86	-8.94	61.15	15.00
21 GLN CA	-19.79	-9.22	60.22	15.00
21 GLN CB	-19.10	-7.92	59.77	15.00
21 GLN CG	-20.08	-6.82	59.31	15.00

-24.20 -11.22

-24.54 -10.58

-22.95 -12.04

-22.93 -12.74

-23.31 -14.09

-24.36 -14.56

**-22.67 -1**.6.23

**-22.36 -16.56** 

-24.46 -16.19

-9.86

-23.44 -17.20 53.45

-24.11 -18.35 54.17

-25.72

58.27

59.46

59.56

58.18

56.78

57.24

55.81

54.34

52.94

56.96

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

25 INH C4

25 INH C5

25 INH C6

25 INH C7

25 INH 08

25 INH C9

25 INH 010

25 INH C11

25 INH C12

25 INH C13

25 INH C14

25 INH C15

PCT/US96/17512

25 INH C16	-21.76	_	56.78	15.00
25 INH 017	-20.87		57.43	15.00
25 INH N18	-21.99	-18.32	56.91	15.00
25 INH C19	-21.15		57.84	15.00
25 INH N20	-22.45	-14.81	56.08	15.00
25 INH C21	-20.81	-20.41	57.54	15.00
25 INH C22	-21.63	-18.99	59.30	15.00
25 INH 023	-21.62	-17.88	59.81	15.00
25 INH C24	-17.90	-23.12	57.09	15.00
25 INH C25	-20.20	-23.31	57.89	15.00
25 INH C26	-21.15	-24.49	58.06	15.00
25 INH C27	-21.28	-25.45	56.87	15.00
25 INH C28	-19.93	-25.80	56.26	15.00
25 INH C29	-22.00	-26.71	57.30	15.00
25 INH C30	-20.37	-22.34	59.06	15.00
25 INH 031	-20.24	-22.76	60.20	15.00
25 INH N32	-20.66	-21.04	58.80	15.00
25 INH C33	-20.82	-20.11	59.89	15.00
25 INH N34	-18.81	-23.74	57.85	15.00
26 TRP N	-22.76	-20.93	62.89	15.00
26 TRP CA	-22.67	-22.24	63.53	15.00
26 TRP CB	-21.35	-22.95	63.19	15.00
26 TRP CG	-20.14	-22.22	63.67	15.00
26 TRP CD2	-19.45	-22.41	64.92	15.00
26 TRP CE2	-18.44	-21.43	65.00	15.00
26 TRP CE3	-19.61	-23.30	65.99	15.00
26 TRP CD1	-19.51	-21.18	63.05	15.00
26 TRP NE1	-18.50	-20.70	63.84	15.00
26 TRP CZ2	-17.59	-21.32	66.10	15.00
26 TRP CZ3	-18.76	-23.19	67.08	15.00
26 TRP CH2	-17.76	-22.21	67.13	15.00
26 TRP C	-22.83	-22.09	65.05	15.00
26 TRP O	-23.50			15.00
27 ALA N	-22.22	-21.05	65. <b>6</b> 0	15.00
27 ALA CA		-20.76	67.03	15.00
27 ALA CB			67.37	15.00
27 ALA C			67.41	15.00
27 ALA O	-24.15		68.53	15.00
28 PHE N	-24.44	<b>-19</b> .79	66.48	15.00
28 PHE CA	25.83	-19.42	65.72	15.00
28 PHE CB			65.81	15.00
28 PHE CG	-25.77	-16.95	66.24	15.00

28 PHE CD		-16.50	65.83	15.00
28 PHE CD		-16.16	67.11	15.00
28 PHE CE	1 -24.03	-15.31	66.28	15.00
28 PHE CE	2 -26.02	-14.96	67.56	15.00
28 PHE CZ	-24.78	-14.53	67.15	15.00
28 PHE C	-26.76	-20.62	66.63	15.00
28 PHE 0	-27.71	-20.74	67.40	15.00
29 SER N	-26.51	-21.50	65.67	15.00
29 SER CA	-27.33	-22.69	65.53	15.00
29 SER CB	-26.88	-23.54	64.35	15.00
29 SER OG	-27.59	-24.76	64.31	15.00
29 SER C	-27.22	-23.49	66.82	15.00
29 SER 0	-28.20	-23.61	67.54	15.00
30 SER N	-26.01	-23.95	67.12	15.00
30 SER CA	-25.70	-24.75	68.32	15.00
30 SER CB	-24.19	-24.86	68.50	15.00
30 SER OG	-23.53	-25.04	67.27	15.00
30 SER C	-26.34	-24.24	69.61	15.00
30 SER O	-26.97	-25.00	70.37	15.00
31 VAL N	-26.15	-22.96	69.88	15.00
31 VAL CA	-26.71	-22.32	71.06	15.00
31 VAL CB	-26.23	-20.85	71.13	15.00
31 VAL CG1		-20.01	71.98	15.00
31 VAL CG2		-20.81	71.69	15.00
31 VAL C		-22.44	71.03	15.00
31 VAL 0	-28.86	-22.74	72.04	15.00
32 GLY N	-28.82	-22.26	69.85	15.00
32 GLY CA	-30.26	-22.36	69.72	15.00
32 GLY C	-30.78	-23.75	70.03	15.00
32 GLY 0	-31.86	-23.89	70.62	15.00
33 ALA N	-30.07	-24.78	69.61	15.00
33 ALA CA	-30.50	-26.14	69.90	15.00
33 ALA CB	-29.65	-27.14	69.16	15.00
33 ALA C		-26.36	71.40	15.00
33 ALA O		-26.85	72.05	15.00
34 LEU N	-29.24	-25.92	71.96	15.00
34 LEU CA	-28.96	-26.04	73.40	15.00
34 LEU CB	-27.57	<b>-25.5</b> 0	73.75	15.00
34 LEU CG	-26.28	-26.23	73.40	15.00
34 LEU CD1		-25.32	73.73	15.00
34 LEU CD2		-27.53	74.19	15.00
34 LEU C	-30.00	-25.35	74.28	15.00

34 LEU O	-30.28	2 . 25 . 01	75.00	
35 GLU N	-30.49			
35 GLU CA	-31.50			
35 GLU CB	-31.65			
35 GLU CG	-30.41		74.01	15.00
35 GLU CD	-30.41		74.08	15.00
35 GLU OE1	-31.23		73.14	
35 GLU 0E2	-29.79			
35 GLU C	-32.84		- <del>-</del>	
35 GLU O	-33.60		74.47	
36 GLY N	-33.60		75.44	15.00
36 GLY CA			73.31	
36 GLY C	-34.38		73.12	15.00
36 GLY O	-34.45		74.11	15.00
37 GLN N	-35.47		74.79	15.00
37 GLN CA	-33.35	-27.32	74.22	15.00
37 GLN CB	-33.25	-28.43	75.15	15.00
37 GLN CB	-32.05	-29.31	74.83	15.00
	-32.27		73.64	15.00
	-33.50		73.81	15.00
37 GLN 0E1	-33.74	-31.63	74.88	15.00
37 GLN NE2	-34.28	-31.20	72.75	15.00
37 GLN C	-33.22	-27.96	76.60	15.00
37 GLN 0	-33.78	-28.61	77.48	15.00
38 LEU N	-32.60	-26.81	76.84	15.00
38 LEU CA	-32.52	-26.23	78.19	15.00
38 LEU CB	-31.72	-24.94	78.20	15.00
38 LEU CG	-31.57	-24.25	79.55	15.00
38 LEU CD1	-30.61	-25.04	80.42	15.00
38 LEU CD2	-31.08	-22.82	79.38	15.00
38 LEU C	-33.94	-26.00	78.70	15.00
38 LEU O	-34.27	-26.33	79.83	15.00
39 LYS N	-34.78	-25.42	77.86	15.00
39 LYS CA	-36.17	-25.16	78.19	15.00
39 LYS CB	-36.85	-24.36	77.08	15.00
39 LYS CG	-38.38	-24.43	77.06	15.00
39 LYS CD		23.€8	78.21	15.00
39 LYS CE		-23.93	78.24	15.00
39 LYS NZ		-23.27	79.40	15.00
39 LYS C		-26.48	78.42	15.00
39 LYS O		-26.59	79.30	15.00
40 LYS N		-27.49	77.63	15.00
40 LYS CA	-37.2C	-28.79	77.78	15.00

45 LEU CG

45 LEU CD1

45 LEU CD2

-35.60

-35.30

PCT/US96/17512

76.34

75.64

75.55

15.00

15.00

15.00

-19.92

-21.23

-36.62 -19.13

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45 LEU C	-34.32	-19.32	79.23	15.00
45 LEU O	-34.65	-18.34	79.89	15.00
46 LEU N	-33.11	-19.46	78.68	15.00
46 LEU CA	-32.02	-18.47	78.83	15.00
46 LEU CB	-30.97	-18.94	79.82	15.00
46 LEU CG	-30.95	-18.53	81.29	15.00
46 LEU CD1	-29.80	-19.23	81.98	15.00
46 LEU CD2	-30.80	-17.02	81.38	15.00
46 LEU C	-31.33	-18.24	77.48	15.00
46 LEU O	-31.36	-19.10	76.62	15.00
47 ASN N	-30.68	-17.09	77.32	15.00
47 ASN CA	-29.95	-16.81	76.10	15.00
47 ASN CB	-29.88	-15.32	75.78	15.00
47 ASN CG	-31.23	-14.72	75.46	15.00
47 ASN OD1	-31.79	-13.96	76.25	15.00
47 ASN ND2	-31.74	-15.03	74.28	15.00
47 ASN C	-28.56	-17.33	76.35	15.00
47 ASN 0	-27.87	-16.81	77.23	15.00
48 LEU N	-28.16	-18.41	75.67	15.00
48 LEU CA	-26.81	-18.95	75.85	15.00
48 LEU CB	-26.75	-20.44	75.51	15.00
48 LEU CG	-27.61	-21.41	76.33	15.00
48 LEU CD1	-27.10	-22.82	76.11	15.00
48 LEU CD2	-27.55	-21.06	77.80	15.00
48 LEU C	-25.82	-18.14	75.02	15.00
48 LEU O	-26.22	-17.32	74.19	15.00
49 SER N	-24.53	-18.38	75.23	15.00
49 SER CA	-23.48	-17.62	74.55	15.00
49 SER CB	-22.43	-17.16	75.56	15.00
49 SER OG	-21.36	-16.51	74.91	15.00
49 SER C	-22.77	-18.19	73.33	15.00
49 SER O	-21.87	-19.03	73.46	15.00
50 PRO N	-23.11	-17.69	72.12	15.00
50 PRO CD	-24.26	-16.83	71.79	15.00
50 PRO CA	-22.46	-18.17	70.90	15.00
50 PRO CB	-23.32	-17.56	69.80	15.00
50 PRC CG	-23.89	-16.34	70.43	15.00
50 PRO C	-21.01	-17.66	70.85	15.00
50 PRO O	-20.16	-18.22	70.16	15.00
51 GLN N	-20.74	-16.59	71.61	15.00
51 GLN CA	-19.41	-15.98	71.71	15.00
51 GLN CB	-19.50	-14.57	72.29	15.00

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51 GLN CG		-13.80	72.34	15.00
51 GLN CD	-17.66	-13.43	70.96	15.00
51 GLN OE1	-18.33	-12.73	70.19	15.00
51 GLN NE2	-16.45	-13.88	70.65	15.00
51 GLN C	-18.50	-16.86	72.56	15.00
51 GLN 0	-17.27	-16.82	72.42	15.00
52 ASN N	-19.11	-17.64	73.44	15.00
52 ASN CA	-18.38	-18.58	74.30	15.00
52 ASN CB	-19.35	-19.26	75.26	15.00
52 ASN CG	-18.67	-20.14	76.30	15.00
52 ASN OD1	-18.91	-19.99	77.51	15.00
52 ASN ND2	-17.88	-21.11	75.85	15.00
52 <b>ASN</b> C	-17.74	-19.57	73.32	15.00
52 ASN 0	-16.55	-19.90	73.41	15.00
53 LEU N	-18.55	-20.00	72.36	15.00
53 LEU CA	-18.11	-20.96	71.35	15.00
53 LEU CB	-19.32	-21.52	70.61	15.00
53 LEU CG	-20.38	-22.09	71.55	15.00
53 LEU CD1	-21.65	-22.44	70.79	15.00
53 LEU CD2	-19.83	-23.30	72.25	15.00
53 LEU C	-17.06	-20.37	70.39	15.00
53 LEU O	-15.94	-20.89	70.28	15.00
54 VAL N	-17.43	-19.27	69.73	15.00
54 VAL CA	-16.55	-18.58	68.78	15.00
54 VAL CB	-17.06	-17.14	68.52	15.00
54 VAL CG1	-16.12	-16.40	67.60	15.00
54 VAL CG2	-18.46	-17.18	67.91	15.00
54 VAL C	-15.12	-18.49	69.29	15.00
54 VAL O	-14.16	-18.68	68.55	15.00
55 ASP N	-15.00	-18.21	70.58	15.00
55 ASP CA	-13.71	-18.08	71.23	15.00
55 ASP CB	-13.82	-17.19	72.49	15.00
55 ASP CG	-14.16	-15.76	72.16	15.00
55 ASP OD1	-13.98	-15.35	71.00	15.00
55 ASP OD2	-14.62	-15.03	73.06	15.00
55 ASP C	-13.05	-19.41	71.60	15.00
55 ASP O	-11.98	-19.76	71.08	15.00
56 CYS N	-13.74	-20.18	72.43	15.00
56 CYS CA	-13.21	-21.42	72.96	15.00
56 CYS C	-13.19	-22.70	72.14	15.00
56 CYS O	-12.40	-23.59	72.45	15.00
56 CYS CB	-13.84	-21.68	74.32	15.00

WO 97/16177		TABLE	X	
56 CYS SG	-14.09	-20.14	75.26	15.00
57 VAL N	-14.06	-22.83	71.15	15.00
57 VAL CA	-14.11	-24.04	70.33	15.00
57 VAL CB	-15.47	-24.21	69.61	15.00
57 VAL CG1	-15.58	-25.61	69.01	15.00
57 VAL CG2	-16.61	-23.97	70.58	15.00
57 VAL C	-12.98	-24.06	69.30	15.00
57 VAL O	-13.18	-23.78	68.12	15.00
58 SER N	-11.80	-24.45	69.76	15.00
58 SER CA	-10.60	-24.55	68.94	15.00
58 SER CB	-9.45	-25.07	69.79	15.00
58 SER OG	-9.53	-24.53	71.10	15.00
58 SER C	-10.73	-25.37	67. <b>6</b> 7	15.00
58 SER O	-9.99	-25.17	66.72	15.00
59 GLU N	-11.61	-26.36	67.70	15.00
59 GLU CA	-11.83	-27.23	66.55	15.00
59 GLU CB	-12.73	-28.41	66.92	15.00
59 GLU CG	-12.20	-29.30	68.03	15.00
59 GLU CD	-12.38	-28.71	69.41	15.00
59 GLU OE1	-13.54	-28.51	69.82	15.00
59 GLU OE2	-11.37	-28.43	70.06	15.00
59 GLU C	-12.41	-26.48	65.37	15.00
59 GLU O	-12.37	-26. <b>9</b> 5	64.23	15.00
60 ASN N	-13.03	-25.34	65.65	15.00
60 ASN CA	-13.65	-24.52	64.62	15.00
60 ASN CB	-15.10	-24.18	64.99	15.00
60 ASN CG	-16.04	-25.37	64.87	15.00
60 ASN OD1	-17.24	-25.24	65.03	15.00
60 ASN ND2	-15.49	-26.53	64.55	15.00
60 ASN C	-12.83	-23.26	64.38	15.00
60 ASN O	-11.82	-23.03	65. <b>05</b>	15.00
61 ASP N	-13.28	-22.43	63.44	15.00
61 ASP CA	-12.56	-21.22	63.09	15.00
61 ASP CB	-12.53	-21.05	61.57	15.00
C1 100 00				

-11.12 -20.82

-10.18 -20.65

-10.96 -20.82

**-13.09 -19.95** 

-12.67 -18.85

-14.00 -20.09

**-15.40 --18.19** 

-18.91

-14.55

61 ASP CG

61 ASP OD1

61 ASP OD2

61 ASP C

61 ASP O

62 GLY N

62 GLY CA

62 GLY C

PCT/US96/17512

61.03

61.83

59.79

63.76

63.43

64.72

65.36

64.33

15.00

15.00

15.00

15.00

15.00

15.00

15.00

62 GLY O	-16.39	-18.75	63.85	15.00
63 CYS N	-14.98	-16.99	63.94	15.00
63 CYS CA	-15.70	-16.19	62.94	15.00
63 CYS C	-15.40	-16.66	61.53	15.00
63 CYS O	-15.88	-16.07	60.57	15.00
63 CYS CB	-15.39	-14.70	63.02	15.00
63 CYS SG	-16.14	-13.86	64.44	15.00
64 GLY N	-14.57	-17.68	61.40	15.00
64 GLY CA	-14.26	-18.20	60.08	15.00
64 GLY C	-15.21	-19.31	59.69	15.00
64 GLY O	-15.23	-19.74	58.53	15.00
65 GLY N	-15.99	-19.80	60.65	15.00
65 GLY CA	-16.93	-20.87	60.37	15.00
65 GLY C	-16.62	-22.14	61.14	15.00
65 GLY O	-15.48	-22.38	61.54	15.00
66 GLY N	-17.64	-22.95	61.35	15.00
66 GLY CA	-17.45		62.08	15.00
66 GLY C	-18.56	-25.20	61.92	15.00
66 GLY O	-19.38	-25.12	61.00	15.00
67 TYR N	-18.56	-26.18	62.81	15.00
67 TYR CA	-19.55	-27.25	62.80	15.00
67 TYR CB	-18.89		62.62	15.00
67 TYR CG	-18.09		61.37	15.00
67 TYR CD1	-18.68	-28.65	60.11	15.00
67 TYR CE1	-17.97	-28.95	58.95	15.00
67 TYR CD2	-16.77	-29.18	61.44	15.00
67 TYR CE2	-16.05	-29.48	60.30	15.00
67 TYR CZ	-16.65	-29.37	59.06	15.00
67 TYR OH	-15.93	-29.71	57. <b>94</b>	15.00
67 TYR C	-20.31	-27.25	64.11	15.00
67 TYR O	-19.73	-26.97	65.15	15.00
68 MET N	-21.60	-27.57	64.06	15.00
68 MET CA	-22.40	-27.64	65.27	15.00
68 MET CB	-23.88	-27.79	64.94	15.00
68 MET CG		-26.62	64.20	15.00
68 MET SD	-23.93		62.51	15.00
68 MET CE		-27.25	61.69	15.00
68 MET C	-21.91	-28.78	66.17	15.00
68 MET O	-21.81	-28.64	67.39	15.00
69 THR N		-29.90	65.54	15.00
69 THR CA			66.27	15.00
69 THR CB	-20.66	-32.20	65.31	15.00

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69 THR OG1	-19.57		64.48	15.00
69 THR CG2	-21.84	-32.58	64.45	15.00
69 THR C	-19.91	-30.70	67.18	15.00
69 THR O	-19.94	-31.00	68.37	15.00
70 ASN N	-18.92	-30.00	66.64	15.00
70 ASN CA	-17.77	-29.54	67.42	15.00
70 ASN CB	-16.75	-28.84	66.55	15.00
70 ASN CG	-15.86	-29.80	65.85	15.00
70 ASN OD1	-15.33	-30.72	66.46	15.00
70 ASN ND2	-15.68	-29.61	64.55	15.00
70 ASN C	-18.17	-28.63	68.60	15.00
70 ASN O	-17.53	-28.66	69.66	15.00
71 ALA N	-19.20	-27.82	68.40	15.00
71 ALA CA	-19.67	-26.91	69.44	15.00
71 ALA CB	-20.66	-25.91	68.86	15.00
71 ALA C	-20.33	-27.72	70.55	15.00
71 ALA O	-20.26	-27.37	71.72	15.00
72 PHE N	-20.96	-28.83	70.16	15.00
72 PHE CA	-21.61	-29.70	71.13	15.00
72 PHE CB	-22.57	-30.66	70.43	15.00
72 PHE CG	-23.73	-29.98	69.79	15.00
72 PHE CD1	-24.28	-28.84	70.36	15.00
72 PHE CD2	-24.29	-30.48	68.63	15.00
72 PHE CE1	-25.35	-28.21	69.79	15.00
72 PHE CE2	-25.37	-29.87	68.04	15.00
72 PHE CZ	-25.91	-28.72	68.62	15.00
72 PHE C	-20.59	-30.46	71.96	15.00
72 PHE 0	-20.79	-30.69	73.15	15.00
73 GLN N	-19.48	-30.82	71.33	15.00
73 GLN CA	-18.43	-31.54	72.03	15.00
73 GLN CB	-17.46	-32.17	71.04	15.00
73 GLN CG	-16.71	-33.36	71.59	15.00
73 GLN CD	-16.83	-34.56	70.67	15.00
73 GLN OE1	-17.35	-35.61	71.07	15.00
73 GLN NE2	-16.37	-34.41	69.44	15.00
73 GLN C	-17.70	-30.62	72.99	15.00
73 GLN 0	-17.18	-31.07	74.02	15.00
74 TYR N	-17.64	-29.34	72.65	15.00
74 TYR CA	-16.96	-28.38	73.52	15.00
74 TYR CB	-16.74		72.81	15.00
74 TYR CG	-16.38		73.78	15.00
74 TYR CD1	-15.16		74.45	15.00

74 TYR CE1	-14.87	-25.02	75.40	15.00
74 TYR CD2	-17.30	-24.94	74.11	15.00
74 TYR CE2	-17.01	-24.00	75.07	15.00
74 TYR CZ	-15.79		75.71	15.00
74 TYR OH	-15.47		76.67	15.00
74 TYR C	-17.69	-28.17	74.84	15.00
74 TYR O	-17.07	-28.14	75.89	15.00
75 VAL N	-19.00	-27.98	74.77	15.00
75 VAL CA	-19.82	-27.78	75.97	15.00
75 VAL CB	-21.29	-27.47	75.58	15.00
75 VAL CG1	-22.13	-27.23	76.82	15.00
75 VAL CG2	-21.34	-26.25	74.67	15.00
75 VAL C	-19.73	-29.01	76.87	15.00
75 VAL O	-19.82	-28.91	78.10	15.00
76 GLN N	-19.47	-30.16	76.26	15.00
76 GLN CA	-19.33	-31.42	76.97	15.00
76 GLN CB	-19.57	-32.58	76.01	15.00
76 GLN CG	-19.54	-33.95	76.62	15.00
76 GLN CD	-19.67	-35.03	75.58	15.00
76 GLN OE1	-20.73	-35.62	75.41	15.00
76 GLN NE2	-18.60	-35.28	74.86	15.00
76 GLN C	-17.96	-31.55	77.68	15.00
76 GLN O	-17.91	-31.76	78.89	15.00
77 LYS N	-16.87	-31.41	76.94	15.00
77 LYS CA	-15.53	-31.51	77.53	15.00
77 LYS CB	-14.43	-31.56	76.46	15.00
77 LYS CG	-14.07	-30.18	75.87	15.00
77 LYS CD	-12.80	-30.21	75.01	15.00
77 LYS CE	-13.01	-30.90	73.67	15.00
77 LYS NZ	-14.16	-30.32	72.90	15.00
77 LYS C	-15.26	-30.36	78.49	15.00
77 LYS O	-14.45	-30.49	79.41	15.00
78 ASN N	-15.89	-29.22	78.22	15.00
78 ASN CA	-15.73		79.05	15.00
78 ASN CB		-26.77	78.27	15.00
78 ASN CG	-15.64	-25.51	79.00	15.00
78 ASN OD1	-14.49	-25.35	79.40	15.00
78 ASN ND2	-16.57	-24.59	79.14	15.00
78 ASN C		-28.15	80.26	15.00
78 ASN O		-27.45	81.25	15.00
79 ARG N		-29.04	80.16	15.00
79 ARG CA	-18.61	-29.26	81.22	15.00

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79 ARG CB	-17.95	-29.71	82.54	15.00
79 ARG CG	-17.19	-31.05	82.52	15.00
79 ARG CD	-18.12	-32.28	82.55	15.00
79 ARG NE	-18.94	-32.34	83.76	15.00
79 ARG CZ	-20.14	-32.92	83.84	15.00
79 ARG NH1	-20.69	-33.51	82.78	15.00
79 ARG NH2	-20.82	-32.88	84.99	15.00
79 ARG C	-19.47	-28.02	81.44	15.00
79 ARG O	-19.86	-27.74	82.57	15.00
80 GLY N	-19.75	-27.27	80.38	15.00
80 GLY CA	-20.58	-26.08	80.52	15.00
80 GLY C	-20.38	-24.97	79.49	15.00
80 GLY O	-19.36	-24.93	78.78	15.00
81 ILE N	-21.37	-24.08	79.41	15.00
81 ILE CA	-21.35	-22.92	78.50	15.00
81 ILE CB	-22.14	-23.20	77.17	15.00
81 ILE CG2	-23.59		77.46	15.00
81 ILE CG1	-22.11	-21.97		15.00
81 ILE CD1	-22.75	-22.19	74.90	15.00
81 ILE C	-22.00	-21.76	79.25	15.00
81 ILE O	-22.86	-21.98	80.11	15.00
82 ASP N	-21.56	-20.54	78.96	15.00
82 ASP CA	-22.09			15.00
82 ASP CB	-21.08	-18.20	79.59	15.00
82 ASP CG	-19.89	-18.44	80.48	15.00
82 ASP OD1	-18.82	-17.87	80.21	15.00
82 ASP OD2	-20.03	-19.18	81.47	15.00
82 ASP C	-23.40	-18.85	79.02	15.00
82 ASP 0	-23.89	-19.36	78.02	15.00
83 SER N	-23.96	-17.84	79.68	15.00
83 SER CA	-25.19	-17.20	79.27	15.00
83 SER CB	-26.03	-16.83	80.49	15.00
83 SER OG	-25.19	-16.35		15.00
83 SER C	-24.76	-15.96	78.49	15.00
83 SER O	-23.59	-15.58	78.53	15.00
84 GLU N	-25.68	-15.29	77.81	15.00
84 GLU CA	-25.29	-14.13	77.03	15.00
84 GLU CB	-26.39	-13.59	76.13	15.00
84 GLU CG		-12.69	75.03	15.00
84 GLU CD		-13.42	74.11	15.00
84 GLU OE1		-13.79	72.99	15.00
84 GLU OE2	-23.68	-13.64	74.49	15.00
			73.37	13.00

84 GLU C	-24.73	-13.05	77.92	15.00
84 GLU O	-23.52	-12.84	77.90	15.00
85 ASP N	-25.56	-12.36	78.69	15.00
85 ASP CA	-25.00	-11.32	79.55	15.00
85 ASP CB	-26.06	-10.32	80.06	15.00
85 ASP CG	-25.45	-8.93	80.42	15.00
85 ASP OD1	-26.14	-8.14	81.10	15.00
85 ASP OD2	-24.30	-8.62	80.02	15.00
85 ASP C	-24.32	-12.06	80.68	15.00
85 ASP O	-24.91	-12.29	81.73	15.00
86 ALA N	-23.11	-12.51	80.37	15.00
86 ALA CA	-22.20	-13.27	81.22	15.00
86 ALA CB	-22.80	-14.62	81.60	15.00
86 ALA C	-20.96	-13.46	80.33	15.00
86 ALA O	-19.83	-13.52	80.81	15.00
87 TYR N	-21.21	-13.59	79.03	15.00
87 TYR CA	-20.18	-13.73	78.00	15.00
87 TYR CB	-19.74	-15.19	77.86	15.00
87 TYR CG	-18.39	-15.40	77.19	15.00
87 TYR CD1	-17.78	-14.38	76.45	15.00
87 TYR CE1	-16.54	-14.59	75.84	15.00
87 TYR CD2	-17.73	-16.61	77.31	15.00
87 TYR CE2	-16.49	-16.82	76.72	15.00
87 TYR CZ	-15.90	-15.80	75 <b>.9</b> 8	15.00
87 TYR OH	-14.67	-15.99	75.42	15.00
87 TYR C	-20.88	-13.21	76.73	15.00
87 TYR O	-21.25	-13.98	75.86	15.00
88 PRO N	-21.07	-11.89	76.64	15.00
88 PRO CD	-20.61	-10.92	77.65	15.00
88 PRO CA	-21.72	-11.18	75.54	15.00
88 PRO CB	-21.69	-9.73	76.00	15.00
88 PRO CG	-21.61	-9.83	77.49	15.00
88 PRO C	-21.11	-11.32	74.15	15.00
88 PRO 0		-11.51	74.00	15.00
89 TYR N	-21.95		73.14	15.00
89 TYR CA	-21.55	-11.21	71.74	15.00
89 TYR CB	-22.75	-11.61	70.87	15.00
89 TYR CG	-22.36	-12.04	69.48	15.00
89 TYR CD1	-21.49	-13.11	69.29	15.00
89 TYR CE1		-13.47	68.02	
89 TYR CD2		-11.36	68.36	15.00
89 TYR CE2	-22.41	-11.72	67.10	15.00

89 TYR CZ	-21.53	-12.77	66.94	15.00
89 TYR OH	-21.09	-13.11	65.69	15.00
89 TYR C	-20.90	-9.91	71.24	15.00
89 TYR O	-21.59	-8.91	71.03	15.00
90 VAL N	-19.58	-9.92	71.08	15.00
90 VAL CA	-18.87	-8.73	70.61	15.00
90 VAL CB	-17.41	-8.68	71.09	15.00
90 VAL CG1	-17.35	-8.79	72.60	15.00
90 VAL CG2	-16.59	-9. <b>7</b> 7	70.43	15.00
90 VAL C	-18.89	-8.68	69.08	15.00
90 VAL O	-18.82	-7.61	68.49	15.00
91 GLY N	-18.98	-9.84	68.45	15.00
91 GLY CA	-19.04	-9.87	67.01	15.00
91 GLY C	-17.71	-10.16	66.36	15.00
91 GLY O	-17.56	-9.93	65.16	15.00
92 GLN N	-16.76	-10.69	67.13	15.00
92 GLN CA	-15.43	-11.01	66.62	15.00
92 GLN CB	-14.62	-9.75	66.38	15.00
92 GLN CG	-14.24	-9.09	67.68	15.00
92 GLN CD	-13.83	-7.66	67.52	15.00
92 GLN OE1	-12.91	-7.19	68.19	15.00
92 GLN NE2	-14.53	-6.93	66.64	15.00
92 GLN C	-14.67	-11.92	67.58	15.00
92 GLN O	-14.91	-11.91	68.79	15.00
93 GLU N	-13.72	-12.66	67.02	15.00
93 GLU CA	-12.89	-13.59	67.78	15.00
93 GLU CB	-11.95	-14.35	66.85	15.00
93 GLU CG	-12.64	-15.06	65. <b>69</b>	15.00
93 GLU CD	-11.68	-15.45	64.57	15.00
93 GLU OE1	-10.56	-15.94	64.87	15.00
93 GLU OE2	-12.03	-15.24	63.39	15.00
93 GLU C	-12.11	-12.86	68.86	15.00
93 GLU O	-11.54	-11.79	68.61	15.00
94 GLU N	-12.08	-13.46	70.05	15.00
94 GLU CA	-11.38	<b>-12.9</b> 3	71.22	15.00
94 GLU CB	-12.31	-12.19	72.16	15.00
94 GLU CG	-12.77	-10.82	71.71	15.00
94 GLU CD	-13.69	-10.18	72.74	15.00
94 GLU OE1	-13.59	-8.95	72.97	15.00
94 GLU OE2		-10.92	73.33	15.00
94 GLU C		-14.12	71.96	15.00
94 GLU O	-11.15	-15.26	71.71	15.00

-13.91

-14.96

-14.79

-15.85

-18.17

-16.48

-18.77

-15.11

-14.28

-16.21

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-17.52

-17.93

-17.61

-18.65

-18.50

-17.90

-18.68

-20.17

-21.01

-21.59

-22.73

-23.27

-18.42

-18.80

-17.74

-17.38

-16.16

-15.66

-14.53

-18.81

**-17.30 -18.62** 

**-16.59 -19.46** 

-15.20 -19.31

-16.27

-17.20 --20.84

-15.50 -21.96

-17.80 -22.53

79.15

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79.54

80.24

79.95

80.59

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98 TYR N

98 TYR CA

98 TYR CB

98 TYR CG

98 TYR CD1

98 TYR CE1

98 TYR CD2

98 TYR CE2

98 TYR CZ

98 TYR OH

98 TYR C

98 TYR 0

99 ASN N

99 ASN CA

99 ASN CB

99 ASN CG

99 ASN OD1

99 ASN ND2

99 ASN C

99 ASN 0

100 PRO N

100 PRO CD

PCT/US96/17512 -

81.62

15.00

-22.56 -31.21

106 LYS NZ

106 LYS C	-25.08		79.17	15.00
106 LYS O	-26.18	-29.04	78.65	15.00
107 CYS N	-24.23	-30.15	78.76	15.00
107 CYS CA	-24.52	-31.01	77.62	
107 CYS CB	-24.00	-30.36	76.33	
107 CYS SG	-24.10	-31.36	74.81	15.00
107 CYS C	-23.81	-32.31	77.94	15.00
107 CYS O	-22.94	-32.34	78.81	15.00
108 ARG N	-24.23	-33.40	77.30	15.00
108 ARG CA	-23.60	-34.70	77.51	15.00
108 ARG CB	-24.27	-35.46	78.66	15.00
108 ARG CG	-25.67	-35.93	78.35	15.00
108 ARG CD	-26.16	-36.95	79.38	15.00
108 ARG NE	-27.36	-37.64	78.90	15.00
108 ARG CZ	-27.34		78.20	15.00
108 ARG NH1	-26.18	-39.37	77.92	15.00
108 ARG NH2	-28.48	-39.29	77.75	15.00
108 ARG C	-23.57	-35.53	76.23	15.00
108 ARG O	-23.71	-36.76	76.27	15.00
109 GLY N	-23.37	-34.87	75.10	15.00
109 GLY CA	-23.30	-35.59	73.85	15.00
109 GLY C	-24.02	-34.85	72. <b>7</b> 5	15.00
109 GLY O	-24.50	-33.73	72.95	15.00
110 TYR N	-24.10	-35.48	71.59	15.00
110 TYR CA	-24.77	-34.92	70.44	15.00
110 TYR CB	-23.91	-33.89	69.74	15.00
110 TYR CG	-22.68	-34.45	69.05	15.00
110 TYR CD1	-21.45	-34.47	69.70	15.00
110 TYR CE1	-20.31	-34.89	69.04	15.00
110 TYR CD2	-22.74	-34.88	67.73	15.00
110 TYR CE2	-21.61	-35.30	67.07	15.00
110 TYR CZ	-20.39	<b>-3</b> 5.30	67.72	15.00
110 TYR OH	-19.24	-35.68	67.05	15.00
110 TYR C	-25.16	-36.02	69.48	15.00
110 TYR O	-24.41	-36.99	69.31	15.00
111 ARG N	-26.32	-35.87	68.86	15.00
111 ARG CA	-26.83	-36.84	67.91	15.00
111 ARG CB	-28.14	-37.43	68.41	15.00
111 ARG CG	-28.01	-38.15	69.74	15.00
111 ARG CD	-28.52	-39.57	69.65	15.00
111 ARG NE	-27.97	-40.28	68.49	15.00
111 ARG CZ	-28.38	-41.48	68.10	15.00

55.18

15.00

-32.09 -37.61

117 ASN N

117 ASN CA	-33.37	-38.18	55.61	15.00
117 ASN CB	-33.13	-39.38	56.52	15.00
117 ASN CG	-34.42	-40.02	57.01	15.00
117 ASN OD1	-35.53	-39.60	56.65	15.00
117 ASN ND2	-34.28	-41.06	57.83	15.00
117 ASN C	-34.32	-37.19	56.27	15.00
117 ASN O	-34.40	-37.10	57.50	15.00
118 GLU N	-35.10	-36.50	55.45	15.00
118 GLU CA	-36.05	-35.52	55.96	15.00
118 GLU CB	-36.81	-34.85	54.83	15.00
118 GLU CG	-36.06	-33.74	54.15	15.00
118 GLU CD	-36.96	-32.95	53.26	15.00
118 GLU OE1	-37.20	-33.39	52.11	15.00
118 GLU OE2	-37.46	-31.91	53.72	15.00
118 GLU C	-37.03	-36.09	56.96	15.00
118 GLU O	-37.48	-35.39	57.88	15.00
119 LYS N	-37.39		56.79	15.00
119 LYS CA	-38.33		57.71	15.00
119 LYS CB	-38.85		57.15	15.00
119 LYS CG	-40.37	-39.34	57.05	15.00
119 LYS CD	-40.95	-38.10	56.34	15.00
119 LYS CE	-42.47	-38.10	56.44	15.00
119 LYS NZ	-43.07	-36.84	55.99	15.00
119 LYS C	-37.71	-38.16	59.09	15.00
119 LYS O	-38.36	-37.86	60.09	15.00
120 ALA N	-36.45	-38.57	59.15	15.00
120 ALA CA	-35.77	-38.75	60.44	15.00
120 ALA CB	-34.45	-39.45	60.27	15.00
120 ALA C	-35.56	-37.40	61.08	15.00
120 ALA O	-35.41	-37.31	62.30	15.00
121 LEU N	-35.52	-36.34	60.26	15.00
121 LEU CA	-35.37	-34.98	60.78	15.00
121 LEU CB	-34.93	-33.98	59.69	15.00
121 LEU CG		-32.55	60.19	15.00
121 LEU CD1	-33.44	-32.47	61.03	15.00
121 LEU CD2	-34.62	-31.58	59.02	15.00
121 LEU C	-36.70	-34.57	61.37	15.00
121 LEU O	-36.77	-34.00	62.45	15.00
122 LYS N		-34.88	60.65	15.00
122 LYS CA		-34.58	61.09	15.00
122 LYS CB		-35.16	60.12	15.00
122 LYS CG	-41.58	-34.91	60.49	15.00

WO 97/16177		TABLE I	x	
122 LYS CD	-42.51	-35.55	59.48	15.00
122 LYS CE	-43.96	-35.15	59.73	15.00
122 LYS NZ	-44.83	-35.53	58.58	15.00
122 LYS C	-39.29	-35.19	62.48	15.00
122 LYS O	-39.68	-34.50	63.42	15.00
123 ARG N	-38.95	-36.47	62.59	15.00
123 ARG CA	-39.04	-37.19	63.86	15.00
123 ARG CB	-38.64	-38.65	63.70	15.00
123 ARG CG	-39.66	-39.49	62.94	15.00
123 ARG CD	-39.69	-40.93	63.45	15.00
123 ARG NE	-38.49	-41.71	63.12	15.00
123 ARG CZ	-38.54	-42.87	62.47	15.00
123 ARG NH1	-39.71	-43.37	62.09	15.00
123 ARG NH2	-37.42	-43.54	62.20	15.00
123 ARG C	-38.19	-36.51	64.93	15.00
123 ARG O	-38.63	-36.34	66.05	15.00
124 ALA N	-36.98	-36.12	64.57	15.00
124 ALA CA	-36.0 <del>9</del>	-35.45	65.50	15.00
124 ALA CB	-34.77	-35.16	64.86	15.00
124 ALA C	-36.70	-34.15	66.00	15.00
124 ALA O	-36.77	-33.93	67.21	15.00
125 VAL N	-37.14	-33.31	65.08	15.00
125 VAL CA	-37.72	-32.03	65.45	15.00
125 VAL CB	-38.13	-31.20	64.21	15.00
125 VAL CG1	-38.87	-29.94	64.63	15.00
125 VAL CG2	-36.90	-30.81	63.41	15.00
125 VAL C	-38.90	-32.21	66.39	15.00
125 VAL O	-39.02	-31.48	67.36	15.00
126 ALA N	-39.75	-33.20	66.13	15.00
126 ALA CA	-40.92	-33.43	66.99	15.00
126 ALA CB	-41.96	-34.25	66.25	15.00
126 ALA C	-40.58	-34.09	68.33	15.00
126 ALA O	-41.09	-33.68	69.38	15.00
127 ARG N	-39.75	-35.12	68.30	15.00
127 ARG CA	-39.38	<b>-35.8</b> 3	69.52	15.00
127 ARG CB	-38.82	-37.21	69.20	15.00
127 ARG CG	-39.74	-38.08	68.36	15.00
127 ARG CD	-39.22	-39.50	68.39	15.00
127 ARG NE		<b>-4</b> 0.36	67.30	15.00
127 ARG CZ	-40.95	-40.55	66.96	15.00
127 ARG NH1	-41.25	<b>-41.3</b> 6	65. <b>96</b>	15.00
127 ARG NH2	-41.92	<b>-39.9</b> 0	67 <b>.59</b>	15.00

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-29.27

-27.93

-28.09

-29.11

-28.80

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-27.54

-27.80

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-27.29

-27.74

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-26.63

-28.26 -26.86

-27.32 -25.81

**-27.05 -26.53** 

-27.93 -27.29

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132 SER CA

132 SER CB

132 SER OG

132 SER C

132 SER 0

133 VAL N

133 VAL CA

133 VAL CB

133 VAL CG1

133 VAL CG2

133 VAL C

133 VAL 0

134 ALA N

134 ALA CA

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O 97/16177		TABLE	TX	
134 ALA CB	-25.65	-27.10	57.62	15.00
134 ALA C	-27.62		56.18	15.00
134 ALA O	-27.92	-27.68	55.69	15.00
135 ILE N	-27.80	-25.44	55.56	15.00
135 ILE CA	-28.36	-25.36	54.21	15.00
135 ILE CB	-29.74	-24.70	54.21	15.00
135 ILE CG2	-30.76	-25.58	54.87	15.00
135 ILE CG1	-29.64	-23.31	54.87	15.00
135 ILE CD1	-30.91	-22.52	54.84	15.00
135 ILE C	-27.51	-24.50	53.29	15.00
135 ILE O	-26.46	-23.99	53.67	15.00
136 ASP N	-28.04	-24.32	52.09	15.00
136 ASP CA	-27.44	-23.50	51.05	15.00
136 ASP CB	-27.56	-24.20	49.70	15.00
136 ASP CG	-27.01	-23.38		
136 ASP OD1	-27.62			15.00
136 ASP OD2	-25.96		48.73	15.00
136 ASP C	-28.17	-22.15	51.05	15.00
136 ASP O	-29.30	-22.07	50.57	15.00
137 ALA N	-27.56		51.62	15.00
137 ALA CA	-28.19		51.68	15.00
137 ALA CB	-28.23		53.12	15.00
137 ALA C	-27.52	-18.76	50.80	15.00
137 ALA O 138 SER N	-27.74	-17.56	50.97	15.00
138 SER N	-26.72 -26.00	-19.22	49.84	15.00
138 SER CB	-24.80			15.00
138 SER OG	-25.20	-20.25	48.33 47.66	15.00 15.00
138 SER C	-26.85	-17.71	47.82	15.00
138 SER O	-26.50			15.00
139 LEU N	-27.96			
139 LEU CA	-28.85	-17.84	46.42	15.00
139 LEU CB		-18.82	46.10	15.00
139 LEU CG		-20.08	45.32	15.00
139 LEU CD1		-20.89	45.13	15.00
139 LEU CD2	-29.04		43.97	15.00
139 LEU C	-29.45	-16.51	46.85	15.00
139 LEU O	-29.82	-16.33	48.00	15.00
140 THR N	-29.61	-15.59	45.90	15.00
140 THR CA	-30.17	-14.28		
140 THR CB	-29.95	-13.25	45.03	15.00

140 THR OG1 -29.88 -13.93 43.77 15.00

49.70

50.77

51.15

15.00

15.00

15.00

**-36.95 -11.28** 

-37.17 -13.65

**-35.94 -12.99** 

145 TYR N

145 TYR CA

145	TYR CB	-36.88	-15.04	51.74	15.00
145	TYR CG	-38.04	-15.65	52.51	15.00
145	TYR CD1	-38.88	-16.58	51.91	15.00
145	TYR CE1	-39.95	-17.13	52.62	15.00
145	TYR CD2	-38.29	-15.28	53.83	15.00
145	TYR CE2	-39.35	-15.82	54.53	15.00
145	TYR CZ	-40.17	-16.74	53.93	15.00
145	TYR OH	-41.23	-17.25	54.64	15.00
145	TYR C	-37.88	-12.76	52.15	15.00
145	TYR O	-37.25	-12.09	52.96	15.00
146	SER N	-39.21	-12.78	52.12	15.00
146	SER CA	-40.01	-12.00	53.04	15.00
146	SER CB	-40.30	-10.61	52.47	15.00
146	SER OG	-40.96	-10.71	51.22	15.00
146	SER C	-41.29	-12.72	53.43	15.00
146	SER O	-41.83	-12.48	54.51	15.00
147	LYS N	-41.76	-13.63	52.58	15.00
147	LYS CA	-42.98	-14.39	52.86	15.00
147	LYS CB	-44.22	-13.50	52.79	15.00
147	LYS CG	-44.61	-13.04	51.39	15.00
147	LYS CD	-45.75	-12.03	51.44	15.00
147	LYS CE	-45.98	-11.35	50.09	15.00
147	LYS NZ	-47.01	-10.28	50.19	15.00
147	LYS C	-43.16	-15.60	51.95	15.00
147	LYS O	-42.49	-15.70	50.92	15.00
148	GLY N	-44.05	-16.50	52.36	15.00
148	GLY CA	-44.34	-17.71	51.60	15.00
148	GLY C	-43.51	-18.90	52.07	15.00
148	GLY O	-42.99	-18.90	53.18	15.00
149	VAL N	-43.43	-19.92	51.22	15.00
149		-42.66	-21.12	51.51	15.00
149	VAL CB	-43.53	-22.39	51.40	15.00
	VAL CG1	-42.69	-23.64	51.56	15.00
149	VAL CG2	-44.62	-22.35	52.45	15.00
149	VAL C	-41.53		50.47	15.00
149	VAL O	-41.75	-21.48	49.30	15.00
150	TYR N	-40.33	-20.80	50.90	15.00
150	TYR CA	-39.19	-20.78	50.01	15.00
150	TYR CB	-37.96	-20.25	50.75	15.00
150	TYR CG	-36.72	-20.12	49.90	15.00
150	TYR CD1	-36.64	-19.13	48.91	15.00
150	TYR CE1	-35.51	-19.02	48.09	15.00

-30.95 -24.11

155 CYS N

43.03

44.26

15.00

15.00

-27.66

-28.86

-29.92

-28.43

-25.34

53.59

53.26

54.34

53.10

53.12

15.00

15.00

15.00

15.00

15.00

-23.80

-24.71

-24.59

-26.15

-22.89

160 LEU CG

160 LEU CD1

160 LEU CD2

160 LEU C

62.01

15.00

-35.20 -27.32

166 ALA CB

172 GLN OE1	-47.54	-19.28	55.40	15.00
172 GLN NE2	-46.46	-19.83	53.52	15.00
172 GLN C	-50.46	-23.45	53.67	15.00
172 GLN 0	-51.49	-24.00	54.07	15.00
173 LYS N	-50.30	-23.07	52.41	15.00
173 LYS CA	-51.38	-23.21	51.46	15.00
173 LYS CB	-51.36	-22.10	50.42	15.00
173 LYS CG	-51.38	-20.70	51.04	15.00
173 LYS CD	-52.68	-20.45	51.78	15.00
173 LYS CE	-53.81	-20.28	50.78	15.00
173 LYS NZ	-53.58	-19.10	49.88	15.00
173 LYS C	-51.15	-24.57	50.86	15.00
173 LYS O	-50.84	-24.70	49.68	15.00
174 GLY N	-51.19	-25.57	51.73	15.00
174 GLY CA	-50.98	-26.94	51.31	15.00
174 GLY C	-49.52	-27.32	51.13	15.00
174 GLY 0	-49.20	-28.49	50.87	15.00
175 ASN N	-48.63	-26.35	51.27	15.00
175 ASN CA	-47.21	-26.62	51.11	15.00
175 ASN CB	-46.45	-25.42	50.56	15.00
175 ASN CG	-47.23	-24.69	49.49	15.00
175 ASN OD1	-48.07	-23.83	49.80	15.00
175 ASN ND2	-46.97	-25.03	48.24	15.00
175 ASN C	-46.60	-27.05	52.42	15.00
175 ASN 0	-46.82	-26.41	53.46	15.00
176 LYS N	-45.87	-28.16	52.38	15.00
176 LYS CA	-45.20	-28.68	53.57	15.00
176 LYS CB	-44.69	-30.10	53.34	15.00
176 LYS CG	-45.74	-31.11	52.86	15.00
176 LYS CD	-46.88	-31.26	53.85	15.00
176 LYS CE	-47.97	-32.21	53.34	15.00
176 LYS NZ	-48.80	-31.66	52.19	15.00
176 LYS C	-44.05	-27.71	53.76	15.00
176 LYS O		-27.08	52.80	15.00
177 HIS N	-43.57	-27.54	54.99	15.00
177 HIS CA	-42.45	-26.64	55.21	15.00
177 HIS CB	-42.88	-25.18	54.99	15.00
177 HIS CG	-43.82	-24.66	56.02	15.00
177 HIS CD2	-43.66	-24.45	57.35	15.00
177 HIS ND1	-45.10	-24.21	55.71	15.00
177 HIS CE1	-45.67	-23.75	56.81	15.00
177 HIS NE2	-44.82	-23.88	57.81	15.00

17	7 HIS	С	-41.71	-26.78	56.53	15.00
17	7 HIS	0	-42.23	-27.35	57.50	15.00
17	B TRP	N	-40.49	-26.24	56.54	15.00
178	3 TRP	CA	-39.60	-26.23	57.69	15.00
178	TRP	CB	-38.19	-26.64	57.31	15.00
178	3 TRP	CG	-37.99	-28.06	56.96	15.00
178	TRP	CD2	-38.07	-29.18	57.84	15.00
178	TRP	CE2	-37.75	-30.33	57.09	15.00
178	3 TRP	CE3	-38.38	-29.33	59.20	15.00
178	TRP	CD1	-37.64	-28.56	55.74	15.00
178	TRP	NE1	-37.49	-29.92	55.81	15.00
178	TRP	CZ2	-37.73	-31.61	57.65	15.00
178	TRP	CZ3	-38.37	-30.60	59.76	15.00
178	TRP	CH2	-38.05	-31.72	58.98	15.00
178	TRP	C	-39.54	-24.81	58.28	15.00
178	TRP	0	-39.17	-23.88	57.58	15.00
179	ILE	N	-39.90	-24.63	59.55	15.00
179	ILE	CA	-39.78	-23.31	60.15	15.00
179	ILE	CB	-40.56	-23.19	61.47	15.00
179	ILE	CG2	-40.42	-21.80	62.04	15.00
179	ILE	CG1	-42.03	~23.55	61.26	15.00
179	ILE	CD1	-42.83	-23.54	62.53	15.00
179	ILE	С	-38.30	-23.20	60.48	15.00
179	ILE	0	-37.76	-24.02	61.22	15.00
180		N	-37.61	-22.23	59.88	15.00
180		CA	-36.18	-22.05	60.12	15.00
180			-35.39	-22.09	58.78	15.00
180			-33.93	-21.75	58.98	15.00
180			-35.49	-23.48	58.17	15.00
180			-35.01	-24.57	59.10	15.00
180	ILE		-35.91	-20.73	60.84	15.00
180	ILE		-36.58	-19.74	60.60	15.00
181	LYS		-34.98	-20.76	61.79	15.00
	LYS			-19.57	62.52	15.00
	LYS			-19.82	64.03	15.00
181			-34.13	-18.62	64.86	15.00
181				-19.01	66.30	15.00
181			-33.59	-17.80	67.13	15.00
181				-18.14	68.55	15.00
181	LYS	С	-33.19	-19.24	62.02	15.00
181				-20.05	62.15	15.00
182	ASN I	N	-33.04	-18.08	61.38	15.00

186 GLU CD

186 GLU OE1

186 GLU OE2

186 GLU C

186 GLU O

187 ASN N

187 ASN CA

-36.84

-37.97

-36.35

-35.77

-36.90

-34.84

-35.06

PCT/US96/17512

-7.96

-8.49

-7.22

-9.12

-9.24

-8.32

-7.47

66.04

66.13

66.93

61.86

61.38

61.36

60.20

15.00

15.00

15.00

15.00

15.00

15.00

15.00

61.03

15.00

-36.77 -12.65

192 GLY N

**-40.17 -21.50** 

**-38.86 -22.76** 

-38.25 -23.48

-36.83 -23.91

**-35.91 -22.73** 

**-34.18 -23.17** 

**-39.67 -26.35** 

**-39.50 -26.46** 

-24.05

-24.66

-25.06

-25.24

-33.94

-39.13

-40.03

-38.87

195 LEU O

196 MET N

196 MET CA

196 MET CB

196 MET CG

196 MET SD

196 MET CE

196 MET C

196 MET O

197 ALA N

197 ALA CA

197 ALA CB

PCT/US96/17512

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

15.00

53.45

54.75

53.98

54.21

54.37

53.22

53.97

52.05

51.54

50.04

52.84

53.63

-38.32 -35.42

-35.33 -33.22

-34.27 -33.16

-35.46 -32.84

**-34.37 -32.30** 

-33.25 -33.34

-32.20 -32.93

46.13

48.27

47.66

49.54

50.35

50.49

51.48

15.00

15.00

15.00

15.00

15.00

15.00

15.00

201 ASN ND2

201 ASN C

201 ASN 0

202 ASN N

202 ASN CA

202 ASN CB

202 ASN CG

PCT/US96/17512

202	ASN	OD1	-32.51	-32.44	52.56	15.00
202	ASN	ND2	-30.94	-33.14	51.13	15.00
202	ASN	C	-33.83	-30.96	49.85	15.00
202	ASN	0	-32.62	-30.77	49.68	15.00
203	ALA	. <b>N</b>	-34.73	-30.00	49.69	15.00
203	ALA	H	-35.60	-30.20	50.07	15.00
203	ALA	CA	-34.39	-28.68	49.18	15.00
203	ALA	CB	-35.57	-27.74	49.24	15.00
203	ALA	С	-33.27	-28.06	50.04	15.00
203	ALA	0	-33.33	-28.03	51.26	15.00
204	CYS	N	-32.23	-27.56	49.35	15.00
204	CYS	CA	-31.11	-26.89	50.02	15.00
204	CYS	С	-30.29	-27.71	51.00	15.00
204	CYS	0	-29.50	-27.14	51.76	15.00
204	CYS	CB	-31.58	-25.60	50.69	15.00
204	CYS	SG	-32.12	-24.29	49.55	15.00
205	GLY	N	-30.43	-29.03	50.98	15.00
205	GLY	CA	-29.68	-29.88	51.90	15.00
205	GLY	С	-30.10	-29.71	53.35	15.00
205	GLY	0	-29.31	<b>-29.9</b> 2	54.27	15.00
206	ILE		-31.37	-29.38	53.54	15.00
206		CA	-31.95	-29.16	54.85	15.00
206	ILE	CB	-33.46	-28.92	54.73	15.00
206	ILE	CG2	-34.13	-30.13	54.12	15.00
206		CG1	-34.06	-28.55	56.10	15.00
206	ILE		-33.76	-27.15	56.54	15.00
206	ILE		-31.71	-30.29	55.85	15.00
206	ILE		-31.47	-30.04	57.03	15.00
207	ALA		-31.79	-31.54	55.38	15.00
207	ALA		-31.58	-32.69	56.25	15.00
207	ALA		-32.72	-33.68	56.08	15.00
207	ALA		-30.25	-33.35	55.94	15.00
	ALA		-30.15	-34.57	55.95	15.00
	ASN		-29.24		55.67	15.00
	ASN			-33.04	55.33	15.00
	ASN			-32.42	54.03	15.00
	ASN			-33.30	52.84	15.00
	ASN			-33.50	52.47	15.00
	ASN			-33.86	52.26	15.00
	ASN			-32.89		15.00
	ASN			<b>-33.5</b> 3		15.00
209	LEU	N	-27.08	-32.00	57.36	15.00

-28.21 -32.91

**-29.45 -32.81** 

-27.50 -33.77

-28.12 -34.59

-27.50 -35.97

-28.01 -37.00

-27.20 -38.28

213 PRO C

213 PRO O

214 LYS N

214 LYS CA

214 LYS CB

214 LYS CG

214 LYS CD

PCT/US96/17512

72.65

72.72

73.38

74.42

74.52

73.53

73.68

15.00

15.00

15.00

15.00

15.00

15.00

15.00

		IABLE	LX.	
214 LYS CE	-27.79		72.93	15.00
214 LYS NZ	-27.00	-40.67	73.21	15.00
214 LYS C	-27.86	-33.84	75. <b>7</b> 0	15.00
214 LYS O	-26.73	-33.45	75. <b>9</b> 8	15.00
215 MET N	-28.90	-33.63	76.49	15.00
215 MET CA	-28.73	-32.91	77.73	15.00
215 MET CB	-29.47	-31.58	77.67	15.00
215 MET CG	-28.62	-30.45	78.17	15.00
215 MET SD	-29.49	-28.92	78.17	15.00
215 MET CE	-30.02	-28.81	79.89	15.00
215 MET C	-29.15	-33.74	78.95	15.00
215 MET OT1	-30.12	-34.53	78.84	15.00
215 MET OT2	-28.49	-33.59	80.01	15.00
216 НОН ОН2	-28.59	-18.05	86.43	15.00
217 HOH OH2	-24.24	-33.32	82.08	15.00
218 НОН ОН2	-30.97	-16.19	65.69	15.00
219 НОН ОН2	-30.10	-20.71	63.47	15.00
220 нон он2	-13.66	-11.12	63.10	15.00
221 нон он2	-9.67	-9.48	64.25	15.00
222 нон он2	-34.55	-23.08	70.24	15.00
223 нон он2	-14.15	-32.13	69.51	15.00
224 НОН ОН2	-11.90	-8.52	62.51	15.00
225 нон он2	-24.25	-30.66	62.17	15.00
226 НОН ОН2	-10.58	-2.52	79.25	15.00
227 НОН ОН2	-14.05	-21.32	67.22	15.00
228 нон он2	-44.68	-30.63	50.04	15.00
229 нон он2	-45.38	-36.05	56.05	15.00
230 нон он2	-39.65	-13.31	65.32	15.00
231 нон он2	-35.12	-36.60	49.29	15.00
232 нон он2	-17.36	-34.13	65.07	15.00
233 нон он2	-30.35	-19.53	65.73	15.00
234 нон он2	-27.89	-19.53	62.51	15.00
235 нон он2	-21.85	-29.55	62.34	15.00
236 нон он2	-30. <b>14</b>	3.73	67.17	15.00
237 НОН ОН2	-40.50	-29.62	80.16	15.00
238 нон он2	-27.85	-23.15	86.33	15.00
239 нон он2	-38.29	<b>~13.9</b> 5	44.87	15.00
240 HOH OH2	-36.58	-24.59	50.05	15.00
241 HOH OH2	-46.68	-34.18	57.37	15.00
242 HOH OH2	-26.77	-6.82	59.79	15.00
243 нон он2	-43.58	-17.40	60.45	15.00
244 нон он2	-23.22	-6.13	61.38	15.00

245	нон 3	OH2	-33.13	-28.30	71.09	15.00
246	5 нон	OH2	-46.57	-25.22	78.97	15.00
247	7 нон	OH2	-14.51	-7.76	88.79	15.00
248	нон 8	OH2	-3.26	-20.73	74.76	15.00
249	НОН	OH2	0.44	-15.91	75.31	15.00
250	нон (	OH2	-19.71	-34.82	58.63	15.00
251	нон	OH2	-34.91	-11.28	53.79	15.00
252	HOH	OH2	-32.46	-28.27	46.13	15.00
253	нон	OH2	-38.20	-15.68	37.93	15.00
254	нон	OH2	-41.44	-34.28	56.30	15.00
255	НОН	OH2	-46.93	-13.62	73.92	15.00
256	НОН	OH2	-32.58	-13.60	60.68	15.00
257	HOH	OH2	-35.46	-6.38	55.50	15.00
258	НОН	OH2	-24.79	-7.91	66.67	15.00
259	HOH	OH2	-32.06	-6.48	63.77	15.00
260	HOH	OH2	-17.19	-5.30	66.67	15.00
261	HOH	OH2	-33.68	-20.47	70.17	15.00
262	HOH	OH2	-13.42	-23.06	78.55	15.00
263	HOH	OH2	-8.54	-20.70	73.58	15.00
264	HOH	OH2	-8.22	-29.32	76.42	15.00
265	HOH	OH2	-25.08	-33.76	60.84	15.00
266	HOH	OH2	-23.92	-37.99	66.66	15.00
267	HOH	OH2	-14.04	-33.08	66.81	15.00
268	HOH	OH2	-12.79	-27.03	71.88	15.00
269	HOH	OH2	-18.55	-42.19	77.34	15.00
270	HOH	OH2	-22.19	-37.43	71.34	15.00
271	HOH	OH2	-3.79	-11.43	71.45	15.00
272	HOH	OH2	-10.91	-19.86	67.02	15.00
273	HOH	OH2	-30.22	-20.12	49.07	15.00
274	HOH	OH2	-25.88	-18.93	42.52	15.00
275	нон	OH2	-36.21	-36.23	51.70	15.00
276	нон	OH2	-20.20	-20.55	47.99	15.00
277	нон	OH2	-38.35	-31.19	41.44	15.00
278	HOH	OH2	-37.29	-30.41	51.12	15.00

Table of the orthogonal three dimensional coordinates in Angstroms and B factors (A<sup>2</sup>) for the cathepsin K complex with inhibitor 1-N-(N-imidazole acetylleucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one.

Residue Atom	x	Y	z	В
1 ALA CB	-8.26	15.35	87.29	15.00
1 ALA C	-6.43	14.73	88.90	15.00
1 ALA O	-6.17	15.27	89.97	15.00
1 ALA N	-8.92	14.74	89.58	15.00
1 ALA CA	-7.91	14.50	88.49	15.00
2 PRO N	-5.47	14.25	88.09	15.00
2 PRO CD	-5.62	13.29	86.98	15.00
2 PRO CA	-4.05	14.45	88.44	15.00
2 PRO CB	-3.32	13.49	87.50	15.00
2 PRO CG	-4.27	13.38	86.30	15.00
2 PRO C	-3.55	15.87	88.27	15.00
2 PRO O	-4.33	16.79	88.21	15.00
3 ASP N	-2.23	16.02	88.20	15.00
3 ASP CA	-1.59	17.30	88.03	15.00
3 ASP CB	-0.07	17.14	88.15	15.00
3 ASP CG	0.45	17.62	89.50	15.00
3 ASP OD1	-0.04	17.07	90.52	15.00
3 ASP OD2	1.29	18.57	89.55	15.00
3 ASP C	-1.90	18.00	86.73	15.00
3 ASP O	-1.71	17.44	85.64	15.00
4 SER N	-2.32	19.26	86.85	15.00
4 SER CA	-2.67	20.16	85.75	15.00
4 SER CB	-3.63	19.49	84.75	15.00
4 SER OG	-4.80	19.03	85.40	15.00
4 SER C	-3.32	21.45	86.30	15.00
4 SER O	-3.83	21.46	87.42	15.00
5 VAL N	-3.30	22.53	85 <b>.53</b>	15.00
5 VAL CA	-3.93	23.80	85.94	15.00
5 VAL CB	-3.00	24.65	86. <b>9</b> 0	15.00
5 VAL CG1	-1.73	25.13	86.17	15.00
5 VAL CG2	-3.76	25.89	87.45	15.00
5 VAL C	-4.21	24.62	84.69	15.00
5 VAL O	-3.43	24.58	83.75	15.00
6 ASP N	-5.38	25.23	84.60	15.00

6 ASP CB	
6 ASP CG	5.00
6 ASP OD1	5.00
6 ASP OD2	5.00
6 ASP C -6.08 27.42 84.03 15 6 ASP O -7.06 27.49 84.72 15 7 TYR N -5.27 28.45 83.88 15 7 TYR CA -5.62 29.77 84.41 15 7 TYR CB -4.36 30.64 84.56 15 7 TYR CG -3.46 30.18 85.71 15 7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CE -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR C -6.71 30.50 83.66 15 7 TYR C -6.71 30.50 83.66 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15	5.00
6 ASP O -7.06 27.49 84.72 15 7 TYR N -5.27 28.45 83.88 15 7 TYR CA -5.62 29.77 84.41 15 7 TYR CB -4.36 30.64 84.56 15 7 TYR CG -3.46 30.18 85.71 15 7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR C -6.71 30.50 83.66 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	5.00
7 TYR N -5.27 28.45 83.88 15 7 TYR CA -5.62 29.77 84.41 15 7 TYR CB -4.36 30.64 84.56 15 7 TYR CG -3.46 30.18 85.71 15 7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	5.00
7 TYR CA -5.62 29.77 84.41 15 7 TYR CB -4.36 30.64 84.56 15 7 TYR CG -3.46 30.18 85.71 15 7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CE -2.00 29.10 87.79 15 7 TYR C -2.00 29.10 87.79 15 7 TYR C -6.71 30.50 83.66 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	5.00
7 TYR CB -4.36 30.64 84.56 15 7 TYR CG -3.46 30.18 85.71 15 7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
7 TYR CG -3.46 30.18 85.71 15 7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
7 TYR CD1 -3.91 30.31 87.05 15 7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
7 TYR CE1 -3.20 29.78 88.07 15 7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
7 TYR CD2 -2.23 29.50 85.46 15 7 TYR CE2 -1.52 28.98 86.45 15 7 TYR CZ -2.00 29.10 87.79 15 7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
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7 TYR OH -1.35 28.46 88.84 15 7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
7 TYR C -6.71 30.50 83.66 15 7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
7 TYR O -7.24 31.48 84.14 15 8 ARG N -7.05 29.98 82.49 15 8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
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8 ARG CA -8.10 30.59 81.68 15 8 ARG CB -8.09 30.02 80.26 15	.00
8 ARG CB -8.09 30.02 80.26 15	.00
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8 ARG CG -6.79 30.28 79.56 15	.00
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10 LYS CG -8.45 23.89 87.51 15	

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10 LYS CD	-7.03	28.46	87.81	15.00
10 LYS CE	-6.43			
10 LYS NZ	-7.24	–		
10 LYS C	-9.98		86.61	
10 LYS O	-10.16			
11 GLY N	-9.85		85.41	
11 GLY CA	-9.94	34.41		
11 GLY C	-8.63	35.17		
11 GLY O	-8.61	36.39		15.00
12 TYR N	-7.54	34.42	85.20	
12 TYR CA	-6.21	34.98	85.26	
12 TYR CB	-5.24	33.95		
12 TYR CG	-5.32			
12 TYR CD1	-6.52		87.99	
12 TYR CE1	-6.58	33.26	89.37	15.00
12 TYR CD2	-4.18	33.66	88.13	15.00
12 TYR CE2	-4.23	33.42	89.47	
12 TYR CZ	-5.42	33.23		15.00
12 TYR OH	-5.43	33.02	91.43	15.00
12 TYR C	-5.68	35.43	83.90	15.00
12 TYR O	-4.68	36.17	83.83	15.00
13 VAL N	-6.34	35.06	82.81	15.00
13 VAL CA	-5.82	35. <b>4</b> 5	81.50	15.00
13 VAL CB	-5.54	34.18	80.66	15.00
13 VAL CG1	-4.93	34.52	79.31	15.00
13 VAL CG2	-4.58	33.24	81.42	15.00
13 VAL C	-6.76	36.37	80.72	15.00
13 VAL O	-7.95	36.17	80.78	15.00
14 THR N	-6.24	37.41	80.08	15.00
14 THR CA	-7.06	38.29	79.27	15.00
14 THR CB	-6.41	39.67	79.12	15.00
14 THR OG1	-5.06	39.53	78.69	
14 THR CG2	-6. <b>4</b> 8	40.43	80.39	15.00
14 THR C	-7.20	37.68	77.89	15.00
14 THR O	-6.41	36.81	77.48	15.00
15 PRO N	-8.14	38.18	77.09	15.00
15 PRO CD	-9.11	39.26	77.34	15.00
15 PRO CA	-8.33	37.64	75.75	15.00
15 PRO CB	-9.51	36.47	75.23	15.00
15 PRO CG	-9.35	39.77	75.97	15.00
15 PRO C	-7.09	37.85	74.89	15.00
15 PRO 0	-6.26	38.73	75.17	15.00
16 VAL N	-6.96	37.00	73.87	15.00

16	VAI	CA	-5.81	37.05	73.00	15.00
16	VAI	CB	-5.74	35.80	72.11	15.00
16	VAI	CG1	-4.42	35.77	71.35	15.00
16	VAI	CG2	-5.84	34.56	72.97	15.00
16	VAL	C	-5.76	38.30	72.12	15.00
16	VAL	. 0	-6.70	38.61	71.37	15.00
17	LYS	N	-4.58	38.91	72.12	15.00
17	LYS	CA	-4.36	40.08	71.32	15.00
17	LYS	CB	-3.75	41.20	72.15	15.00
17	LYS	CG	-4.73	42.01	72.95	15.00
17	LYS	CD	-5.27	41.23	74.12	15.00
17	LYS	CE	-5.71	42.11	75.23	15.00
17	LYS	NZ	-5.93	41.40	76.53	15.00
17	LYS	C	-3.45	39.72	70.16	15.00
17	LYS	0	-2.83	38.63	70.16	15.00
18			-3.32	40.68	69.24	15.00
18	asn	CA	-2.51	40.54	68.03	15.00
18	ASN	CB	-3.37	40.84	66.84	15.00
18			-2.76	40.33	65.56	15.00
18		OD1	-1.57	40.56	65.29	15.00
18	asn	ND2	-3.55	39.60	64.78	15.00
18			-1.33	41.51	68.00	15.00
18			-1.50	42.73	67.86	15.00
19			-0.15	40.93	68.03	15.00
19	GLN		1.12	41.67	68.01	15.00
19	GLN		2.23	40.66	67.93	15.00
19	GLN		3.64	41.21	67.96	15.00
19	GLN		4.65	40.13	68.03	15.00
19		OE1	4.35	39.00	68.42	15.00
19		NE2	5.86	40.47	67.64	15.00
19	GLN		1.22	42.69	66.88	15.00
19	GLN		1.75	43.75	67.04	15.00
20	GLY		0.66	42.38	65.74	15.00
	GLY		0.66	43.27	64.59	15.00
	GLY		1.84	42.90	63.73	15.00
	GLY		1.98	41.74	63.33	15.00
	GLN		2.67	43.89	63.43	15.00
	GLN		3.88	43.70	62.61	15.00
21	GLN		3.90	44.62	61.39	15.00
21			2.90	44.23	60.27	15.00
21	GLN		3.33	43.03	59.44	15.00
	GLN		4.25	43.10	58.59	15.00
21	GLN	NE2	2.58	41.95	59 <b>.5</b> 8	15.00

				А	
	1 GLN C	5.05	44.09	63.52	15.00
2:	_	6.20	43.88	63.19	15.00
22		4.74	44.77	64.60	15.00
22		5.74	45.22	65.53	15.00
22		6.26	44.01	66.35	15.00
22	CYS O	5.48	43.07	66.60	15.00
22	CYS CB	5.08	46.25	66.43	15.00
22	CYS SG	6.07	46.62	67.91	15.00
23	GLY N	7.57	43.99	66.68	15.00
23	GLY CA	8.16	42.91	67.47	15.00
23	GLY C	8.07	43.16	68.96	15.00
23	GLY O	9.08	43.15	69.68	15.00
24	SER N	6.82	43.29	69.42	15.00
24	SER CA	6.47	43.55	70.79	15.00
	SER CB	5.45	44.66	70.86	15.00
24	SER OG	4.31	44.30	70.15	15.00
24	SER C	5.97	42.28	71.46	15.00
24		5.26	42.34	72.45	15.00
25		6.43	41.13	70.98	15.00
25		6.06	39.87	71.53	15.00
25		6.61	38.78	70.65	15.00
25		8.41	38.83	70.46	15.00
25	CYS C	6.49	39.79	73.01	15.00
25		5.74	39.28	73.84	15.00
25		3.24	39.22	63.40	15.00
25		2.86	38.72	62.13	15.00
25		1.57	38.27	61.89	15.00
25	INH C4	0.62	38.31	62.90	15.00
25	INH C5	0.94	38.79	64.16	15.00
25	INH C6	2.25	39.24	64.42	15.00
25	INH O7	4.57	39.75	63.63	15.00
25	INH C8	5.72	39.02	63.91	15.00
25	INH C9	5.62	38.42	65.17	15.00
25		6.58	37.50	65.60	15.00
25	INH C11	7.65	37.17	64.78	15.00
25	INH C12	7.79	37.78	63.51	15.00
25	INH C13	6.82	38.71	63.08	15.00
25	INH S14	8.67	35.93	65.55	15.00
25	INH 015	7.93	34.70	65.54	15.00
25	INH 016	9.92	35.97	64.82	15.00
25	INH N17	8.95	36.39	67.18	15.00
25	INH C18	9.50	37.70	67.57	15.00
25	INH C19	9.05	38.78	68.64	15.00

25	INH	020	8.53	39.68	67.70	15.00
25	INH	C21	10.52	39.34	69.16	15.00
25	INH	N22	11.16	38.50	70.20	15.00
25	INH	C23	12.14	38.91	71.00	15.00
25	INH	024	12.59	40.07	70.97	15.00
25	INH	C25	12.61	37.92	72.07	15.00
25	INH	C26		38.19	73.25	15.00
25	INH	C27	11.80	37.33	74.48	15.00
25	INH	C28	12.06	35.90	74.03	15.00
25	INH	C29	12.95	37.84	75.32	15.00
25	INH	N30	14.03	38.09	72.47	15.00
25	INH	C31	14.92	37.10	72.44	15.00
25	INH	032	14.63	35.96	72.06	15.00
25	INH	C33	16.36	37.38	72.94	15.00
25	INH	C34	17.21	36.17	73.26	15.00
25	INH	C35	17.54	35.58	74.44	15.00
25	INH	N36	18.35	34.51	74.16	15.00
25	INH	C37	18.52	34.43	72.85	15.00
25	INH	N38	17.85	35.42	72.28	15.00
26	TRP	N	7.57	40.50	73.34	15.00
26	TRP	CA	8.08	40.58	74.69	15.00
26	TRP	CB	9.55	41.06	74.66	15.00
26	TRP	CG	9.72	42.45	74.12	15.00
26	TRP	CD2	9.74	43.67	74.85	15.00
26	TRP	CE2	9.80	44.74	73.89	15.00
26	TRP	CE3	9.72	43.99	76.22	15.00
26	TRP	CD1	9.78	42.79	72.83	15.00
26	TRP	NE1	9.82	44.15	72.67	15.00
26	TRP	CZ2	9.83	46.10	74.25	15.00
26	TRP	CZ3	9.75	45.31	76.58	15.00
26	TRP	CH2	9.81	46.37	75.59	15.00
26	TRP	С	7.21	41.51	75.60	15.00
26	TRP	0	7.08	41.27	76.81	15.00
27	ALA	N	6.58	42.54	75.01	15.00
27	ALA	CA	5.77	43.48	75.79	15.00
27	ALA	CB	5.51	44.71	74.96	15.00
27	ALA	С	4.47	42.75	76.05	15.00
27	ALA.	0	3.88	42.85	77.12	15.00
28	PHE	N	4.07	41.90	75.13	15.00
28	PHE	CA	2.81	41.20	75.33	15.00
28	PHE	CB	2.29	40.66	73.99	15.00
28	PHE	CG	1.55	41.63	73.15	15.00
28	PHE	CD1	2.23	42.39	72.20	15.00

28 PHE CD2	0.19	41.92	73.31	15.00
28 PHE CE1	1.57	43.34	71.43	15.00
28 PHE CE2	-0.46	42.87	72.55	15.00
28 PHE CZ	0.23	43.58	71.61	15.00
28 PHE C	2.87	40.12	76.46	15.00
28 PHE O	1.92	39.99	77.26	15.00
29 SER N	3.97	39.39	76.54	15.00
29 SER CA	4.16	38.37	77.55	15.00
29 SER CB	5.40	37.55	77.18	15.00
29 SER OG	5.72	36.66	78.21	15.00
29 SER C	4.28	38.96	78.94	15.00
29 SER O	3.68	38.47	79.91	15.00
30 SER N	5.04	40.05	79.02	15.00
30 SER CA	5.25	40.76	80.28	15.00
30 SER CB	6.13	41.95	80.06	15.00
30 SER OG	7.38	41.52	79.59	15.00
30 SER C	3.96	41.22	80.96	15.00
30 SER O	3.72	40.99	82.17	15.00
31 VAL N	3.13	41.85	80.14	15.00
31 VAL CA	1.83	42.35	80.51	15.00
31 VAL CB	1.33	43.26	79.35	15.00
31 VAL CG1	-0.15	43.27	79.23	15.00
31 VAL CG2	1.86	44.66	79.52	15.00
31 VAL C	0.91	41.16	80.90	15.00
31 VAL O	0.02	41.32	81.77	15.00
32 GLY N	1.16	39.98	80.31	15.00
32 GLY CA	0.35	38.81	80.61	15.00
32 GLY C	0.70	38.25	81.97	15.00
32 GLY O	-0.17	37.79	82.70	15.00
33 ALA N	1.98	38.30	82.34	15.00
33 ALA CA	2.47	37.84	83.63	15.00
33 ALA CB	3.98	37.93	83.67	15.00
33 ALA C	1.84	38.86	84.60	15.00
33 ALA O	1.09	38.48	85.52	15.00
34 LEU N	2.09	40.15	84.37	15.00
34 LEU CA	1.54	41.22	85.20	15.00
34 LEU CB	1.88	42.56	84.56	15.00
34 LEU CG	3.30	43.17	84.66	15.00
34 LEU CD1	3.26	44.56	84.13	15.00
34 LEU CD2	3.75	43.24	86.11	15.00
34 LEU C	0.02	41.13	85.45	15.00
34 LEU O	-0.47	41.28	86.60	15.00
35 GLU N	-0.70	40.80	84.39	15.00

35 GLU CA	-2.16	40.62	84.49	15.00
35 GLU CB	-2.77	40.37	83.11	15.00
35 GLU CG	-2.92	41.61	82.27	15.00
35 GLU CD	-3.06	41.33	80.78	15.00
35 GLU OE1	-2.97	40.14	80.38	15.00
35 GLU OE2	-3.31	42.29	80.00	15.00
35 GLU C	-2.60	39.51	85.47	15.00
35 GLU O	-3.31	39.78	86.45	15.00
36 GLY N	-2.11	38.29	85.24	15.00
36 GLY CA	-2.42	37.14	86.06	15.00
36 GLY C	-2.20	37.38	87.56	15.00
36 GLY O	-2.96	36.89	88.40	15.00
37 GLN N	-1.13	38.09	87.89	15.00
37 GLN CA	-0.78	38.39	89.27	15.00
37 GLN CB	0.64	38.96	89.35	15.00
37 GLN CG	1.72	37.92	89.12	15.00
37 GLN CD	1.50	36.73	90.00	15.00
37 GLN OE1	1.69	36.81	91.20	15.00
37 GLN NE2	1.12	35.61	89.42	15.00
37 GLN C	-1.78	39.38	89.83	15.00
37 GLN O	-2.30	39.19	90.94	15.00
38 LEU N	-2.10	40.38	89.02	15.00
38 LEU CA	-3.04	41.41	89.38	15.00
38 LEU CB	-3.28	42.29	88.15	15.00
38 LEU CG	-4.10	43.56	88.33	15.00
38 LEU CD1	-3.72	44.25	89.66	15.00
38 LEU CD2	-3.95	44.48	87.16	15.00
38 LEU C	-4.32	40.75	89.82	15.00
38 LEU O	-4.90	41.05	90.86	15.00
39 LYS N	-4.75	39.78	89.04	15.00
39 LYS CA	-5.95	39.07	89.36	15.00
39 LYS CB	-6.29	38.15	88.20	15.00
39 LYS CG	-7.34	37.13	88.56	15.00
39 LYS CD	-8.65	37.80	88.80	15.00
39 LYS CE	-9.71	36.74	88.86	15.00
39 LYS NZ	-10.82	37.33	89.61	15.00
39 LYS C	-5.78	38.28	90.67	15.00
39 LYS O	-6.62	38.34	91.53	15.00
10 LYS N	-4.66	37.59	90.83	15.00
0 LYS CA	-4.40	36.83	92.03	15.00
10 LYS CB	-3.01	36.20	91.96	15.00
10 LYS CG	-2.68	35.31	93.16	15.00
10 LYS CD	-1.38	34.53	93.00	15.00

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40 LYS CE	-1.10	33.63	94.23	15.00
40 LYS NZ	-0.27	32.41	93.89	15.00
40 LYS C	-4.50	37.71	93.27	15.00
40 LYS O	-5.26	37.42	94.20	15.00
41 LYS N	-3.86	38.87		15.00
41 LYS CA	-3.77	39.84	94.28	15.00
41 LYS CB	-2.31	40.32	94.41	15.00
41 LYS CG	-1.28	39.21	94.20	15.00
41 LYS CD	-0.04	39.35	95.07	15.00
41 LYS CE	0.68	38.01	95.15	15.00
41 LYS NZ	-0.14	36.86	95.66	15.00
41 LYS C	-4.64	41.07	94.09	15.00
41 LYS O	-4.15	42.18	94.26	15.00
42 THR N	-5.91	40.88	93.78	15.00
42 THR CA	-6.80	42.01	93.60	15.00
42 THR CB	-6.50	42.75	92.28	15.00
42 THR OG1	-5.17	43.26	92.28	15.00
42 THR CG2	-7.48	43.85	92.07	15.00
42 THR C	-8.22	41.47	93.50	15.00
42 THR O	-9.17	42.02	94.07	15.00
43 GLY N	-8.37	40.40	92.74	15.00
43 GLY CA	-9.67	39.80	92.56	15.00
43 GLY C	-10.28	40.20	91.22	15.00
43 GLY O	-11.22	39.57	90.74	15.00
44 LYS N	-9.65	41.19	90.58	15.00
44 LYS CA	-10.13	41.70	89.32	15.00
44 LYS CB	-10.71	43.10	89.52	15.00
44 LYS CG	-11.95	43.16	90.39	15.00
44 LYS CD	-12.36	44.57	90.76	15.00
44 LYS CE	-11.43	45.21	91.78	15.00
44 LYS NZ	-11.50	44.47	93.08	15.00
44 LYS C	-9.02	41.76	88.30	15.00
44 LYS 0	-7.91	42.20	88.61	15.00
45 LEU N	-9.35	41.40	87.07	15.00
45 LEU CA	-8.38	41.39	85.99	15.00
45 LEU CB	-8.61	40.14	85.13	15.00
45 LEU CG	-7.85	39.77	83.85	15.00
45 LEU CD1	-6.47	39.23	84.16	15.00
45 LEU CD2	-8.67	38.72	83.14	15.00
45 LEU C	-8.55	42.68	85.16	15.00
45 LEU O	-9.55	43.36	85.27	15.00
46 LEU N	-7.57	43.04	84.37	15.00
46 LEU CA	-7.65	44.23	83.55	15.00

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46 LEU	CB	-7.27	45.46	84.35	15.00
46 LEU	CG	-7.82	46.76	83.83	15.00
46 LEU	CD1	-9.28	46.67	83.92	15.00
46 LEU	CD2	-7.33	47.91	84.71	15.00
46 LEU	С	-6.63	44.03	82.45	15.00
46 LEU	0	-5.67	43.30	82.61	15.00
47 ASN	N	-6.80	44.76	81.36	15.00
47 ASN	CA	-5.90	44.66	80.25	15.00
47 ASN	CB	-6.61	45.08	78.95	15.00
47 ASN	CG	-7.47	43.98	78.39	15.00
47 ASN	OD1	-7.52	42.88	78.93	15.00
47 ASN	ND2	-8.19	44.28	77.33	15.00
47 ASN	С	-4.77	45.63	80.52	15.00
47 ASN	0	-5.02	46.82	80.64	15.00
48 LEU	N	-3.54	45.17	80.66	15.00
48 LEU	CA	-2.44	46.13	80.87	15.00
48 LEU (	CB	-1.29	45.54	81.75	15.00
48 LEU (		-1.76	45.23	83.19	15.00
48 LEU (		-0.62	44.99	84.10	15.00
48 LEU (		-2.69	46.29	83.75	15.00
48 LEU (		-1.96	46.66	79.51	15.00
48 LEU (		-2.40	46.14	78.46	15.00
49 SER 1		-1.12	47.70	79.52	15.00
49 SER (		-0.63	48.29	78.28	15.00
49 SER (		-0.70	49.82	78.41	15.00
49 SER (		0.06	50.46	77.43	15.00
49 SER (		0.75	47.87	77.84	15.00
49 SER (		1.75	48.29	78.43	15.00
50 PRO N		0.81	47.08	76.75	15.00
50 PRO C		-0.34	46.48	76.03	15.00
50 PRO C		2.10	46.61	76.22	15.00
50 PRO C		1.68	45.45	75.34	15.00
50 PRO C		0.33	45.96	74.79	15.00
50 PRO C				75.45	15.00
50 PRO C		3.99	47.96	75.47	15.00
51 GLN N		1.96	48.67	74.86	15.00
51 GLN C		2.48	49.87	74.18	15.00
51 GLN C		1.37	50.77	73.66	15.00
51 GLN C		2.00	51.92	72.85	15.00
51 GLN C			51.51		15.00
51 GLN O			50.49		15.00
51 GLN N					15.00
51 GLN C	!	3.29	50.70	75 . 17	15.00

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51 GLN O	4.21	51.38	74.79	15.00
52 ASN N	2.93	50.65	76.45	
52 ASN CA	3.62		77.47	15.00
52 ASN CB	2.96	51.15	78.82	
52 ASN CG	3.52	52.04		
52 ASN OD1	4.31	52.93	79.64	
52 ASN ND2	3.09		81.14	
52 ASN C	5.04	50.91	77.52	
52 ASN O	5.98	51.71		15.00
53 LEU N	5.19	49.60	77.57	15.00
53 LEU CA	6.53	49.05	77.64	15.00
53 LEU CB	6.40	47.59	77.98	
53 LEU CG	5.80	47.51	79.34	
53 LEU CD1		46.07	79.79	
53 LEU CD2	6.65	48.34	80.33	15.00
53 LEU C	7.25			15.00
53 LEU O	8.43	49.52	76.27	
54 VAL N	6.53	49.17	75.19	
54 VAL CA	7.19	49.34	73.89	15.00
54 VAL CB	6.25	49.16	72.69	15.00
54 VAL CG1	6.98	49.50	71.41	15.00
54 VAL CG2	5.83	47.75	72.63	15.00
54 VAL C	7.89	50.66		15.00
54 VAL O		50.71	73.32	15.00
55 ASP N		51.70	74.27	15.00
55 ASP CA	7.87	53.05	74.15	15.00
55 ASP CB	6.76	54.08	73.93	15.00
55 ASP CG	5.70	53.58	72.94	15.00
55 ASP OD1		52.76	72.00	15.00
55 ASP OD2		54.09	73.02	15.00
55 ASP C	8.68	53.42	75.33	15.00
55 ASP O	9.58	54.23		15.00
56 CYS N	8.38	52.90	76.50	15.00
56 CYS CA	9.13	53.37	77.67	15.00
56 CYS C	10.33	52.59	78.21	15.00
56 CYS O	11.25	53.15	78.82	15.00
56 CYS CB	8.16	53.80	78.81	15.00
56 CYS SG	6.73	54.73	78.24	15.00
57 VAL N	10.39	51.31	77.97	15.00
57 VAL CA	11.49	50.47	78.46	15.00
57 VAL CB	11.10	48.99	78.38	15.00
57 VAL CG1	12.15	48.16	78.95	15.00
57 VAL CG2	9.83	48.77	79.11	15.00

57 VAL C	12.66		77.55	15.00
57 VAL O	12.94	49.87	76.66	15.00
58 SER N	13.40	51.73	77.79	15.00
58 SER CA	14.54	52.04	76.91	15.00
58 SER CB	15.13	53.44	77.26	15.00
58 SER OG	15.02	53.68	78.65	15.00
58 SER C	15.61	50.98	76.86	15.00
58 SER O	16.39	50.92	75.91	15.00
59 GLU N	15.66	50.15	77.91	15.00
59 GLU CA	16.66	49.07	78.00	15.00
59 GLU CB	16.59	48.34	79.35	15.00
59 GLU CG	16.98	49.19	80.58	15.00
59 GLU CD	15.98	50.33	80.88	15.00
59 GLU OE1	14.75	50.10	80.81	15.00
59 GLU OE2	16.46	51.46	81.17	15.00
59 GLU C	16.48	48.04	76.89	15.00
59 GLU O	17.36	47.20	76.67	15.00
60 asn n	15.31	48.07	76.27	15.00
60 ASN CA	15.00	47.16	75.19	15.00
60 ASN CB	13.64	46.51	75.47	15.00
60 ASN CG	13.69	45.49	76.59	15.00
60 ASN OD1	12.76	45.39	77.36	15.00
60 ASN ND2	14.76	44.68	76.63	15.00
60 ASN C	14.98	47.89	73.85	15.00
60 ASN O	15.26	49.09	73.79	15.00
61 ASP N	14.67	47.16	72.79	15.00
61 ASP CA	14.64	47.72	71.44	15.00
61 ASP CB	15.38	46.80	70.48	15.00
61 ASP CG	16.10	47.57	69.36	15.00
61 ASP OD1	16.04	48.81	69.34	15.00
61 ASP OD2	16.76	46.92	68.52	15.00
61 ASP C	13.28	48.07	70.81	15.00
61 ASP O	13.13	48.13	69.60	15.00
62 GLY N	12.31	48.40	71.63	15.00
62 GLY CA	10.98	48.67	71.11	15.00
62 GLY C	10.49	47.51	70.24	15.00
62 GLY O	10.64	46.38	70.57	15.00
63 CYS N	9.98	47.80	69.05	15.00
63 CYS CA	9.47	46.79	68.11	15.00
63 CYS C	10.61	45.95	67.62	15.00
63 CYS O	10.41	45.05	66.81	15.00
63 CYS CB	8.71	47.45	66.95	15.00
63 CYS SG	7.16	48.19	67.39	15.00

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64 GLY N	11.82	46.25	68.06	15.00
64 GLY CA	12.94	45.47		15.00
64 GLY C	13.32	44.36	68.55	15.00
64 GLY O	14.37	43.74		15.00
65 GLY N	12.48	44.12	69.55	15.00
65 GLY CA	12.76	43.06	70.49	
65 GLY C	13.11	43.58	71.88	
65 GLY O	13.33	44.80	72.13	
66 GLY N	13.21	42.63	72.80	15.00
66 GLY CA	13.50	43.00	74.18	15.00
66 GLY C	13.34	41.82	75.15	15.00
66 GLY O	12.91	40.73	74.72	15.00
67 TYR N	13.65	42.02	76.42	15.00
67 TYR CA	13.55	40.94	77.37	15.00
67 TYR CB	14.85	40.83	78.19	15.00
67 TYR CG	16.13	40.41	77.42	15.00
67 TYR CD1	16.31	39.16	76.96	15.00
67 TYR CE1	17.51	38.78	76.29	15.00
67 TYR CD2	17.13	41.25	77.20	15.00
67 TYR CE2	18.32	40.83	76.53	15.00
67 TYR CZ	18.49	39.61	76.08	15.00
67 TYR OH	19.63	39.12	75.47	15.00
67 TYR C	12.41	41.24	78.31	15.00
67 TYR O	12.00	42.40	78.46	15.00
68 MET N	11.84	40.20	78.88	15.00
68 MET CA	10.72	40.40	79.78	15.00
68 MET CB	10.01	39.09	80.00	15.00
68 MET CG	9.14	38.63	78.85	15.00
68 MET SD	10.15	37.92	77.61	15.00
68 MET CE	10.37	36.10	78.19	15.00
68 MET C	11.20	41.05	81.10	15.00
68 MET O	10.55	41.92	81.68	15.00
69 THR N	12.33	40.56	81.59	15.00
69 THR CA	12.91	41.02	82.85	15.00
69 THR CB	14.24	40.26	83.21	15.00
69 THR OG1	15.16	40.38	82.12	15.00
59 THR CG2	13.99	38.77	83.56	15.00
59 THR C	13.15	42.55	82.80	15.00
59 THR O	13.15	43.20	83.83	15.00
70 ASN N	13.42	43.10	81.63	15.00
0 ASN CA	13.61	44.55	81.51	15.00
0 ASN CB	14.23	44.95	80.15	15.00
0 ASN CG	15.73	44.64	8008	15.00

70 ASN OD1	16.28	44.44	78.98	15.00
70 ASN ND2	16.41	44.61	81.22	15.00
70 ASN C	12.27	45.27	81.66	15.00
70 ASN 0	12.24	46.42	82.08	15.00
71 ALA N	11.20	44.62	81.19	15.00
71 ALA CA	9.84	45.13	81.24	15.00
71 ALA CB	8.96	44.17	80.50	15.00
71 ALA C	9.39	45.23	82.69	15.00
71 ALA O	8.79	46.23	83.13	15.00
72 PHE N	9.72	44.17	83.43	15.00
72 PHE CA	9.39	44.09	84.87	15.00
72 PHE CB	9.67	42.66	85.39	15.00
72 PHE CG	8.80	41.59	84.75	15.00
72 PHE CD1	7.52	41.87	84.33	15.00
72 PHE CD2	9.28	40.32	84.57	15.00
72 PHE CE1	6.73	40.91	83.73	15.00
72 PHE CE2	8.50	39.35	83.98	15.00
72 PHE CZ	7.23	39.64	83.56	15.00
72 PHE C	10.19	45.09	85.71	15.00
72 PHE 0	9.73	45.60	86.72	15.00
73 GLN N	11.41	45.38	85.28	15.00
73 GLN CA	12.27	46.36	85.97	15.00
73 GLN CB	13.74	46.30	85.47	15.00
73 GLN CG	14.74	46.94	86.38	15.00
73 GLN CD	14.58	46.42	87.78	15.00
73 GLN OE1	14.18	45.26	87.96	15.00
73 GLN NE2	14.76	47.28	88.79	15.00
73 GLN C	11.74	47.79	85.84	15.00
73 GLN O	11.83	48.57	86.78	15.00
74 TYR N	11.23	48.14	84.67	15.00
74 TYR CA	10.67	49.45	84.41	15.00
74 TYR CB	10.48	49.62	82.93	15.00
74 TYR CG	9.52	50.72	82.59	15.00
74 TYR CD1	9.90	51.97	82.44	15.00
74 TYR CE1	8.98	53.00	82.06	15.00
74 TYR CD2	8.25	50.50	82.38	15.00
4 TYR CE2	7.39	51.55	82.01	15.00
4 TYR CZ	7.74	52.77	81.85	15.00
4 TYR OH	6.87	53.67	81.39	15.00
4 TYR C	9.35	49.64	85.16	15.00
4 TYR O	9.04	50.74	85.58	15.00
5 VAL N	8.60	48.58	85.39	15.00
5 VAL CA	7.35	48.70	86.13	15.00

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75 VAL CB	6.46	47.47	85.92	15.00
75 VAL CG1	5.15	47.64	86.74	15.00
75 VAL CG2	6.10	47.32	84.44	15.00
75 VAL C	7.63	48.95	87.62	15.00
75 VAL O	6.92	49.76	88.26	15.00
76 GLN N	8.69	48.31	88.13	15.00
76 GLN CA	9.12	48.47	89.51	15.00
76 GLN CB	10.10	47.35	89.89	15.00
76 GLN CG	10.55	47.31	91.38	15.00
76 GLN CD	11.76	46.41	91.65	15.00
76 GLN OE1	12.14	46.22	92.80	15.00
76 GLN NE2	12.37	45.85	90.59	15.00
76 GLN C	9.78	49.88	89.59	15.00
76 GLN O	9.37	50.72	90.39	15.00
77 LYS N	10.72	50.19	88.69	15.00
77 LYS CA	11.40	51.49	88.72	15.00
77 LYS CB	12.56	51.50	87.72	15.00
77 LYS CG	13.20	52.85	87.44	15.00
77 LYS CD	14.22	52.70	86.33	15.00
77 LYS CE	15.18	51.56	86.62	15.00
77 LYS NZ	16.20	51.34	85.56	15.00
77 LYS C	10.51	52.71	88.55	15.00
77 LYS O	10.73	53.76	89.17	15.00
78 ASN N	9.50	52.55	87.70	15.00
78 ASN CA	8.51	53.59	87.37	15.00
78 ASN CB	7.84	53.25	86.04	15.00
78 ASN CG	7.15	54.42	85.42	15.00
78 ASN OD1	7.70	55.49	85.36	15.00
78 ASN ND2	5.93	54.22	84.95	15.00
78 ASN C	7.44	53.69	88.42	15.00
78 ASN O	6.89	54.77	88.64	15.00
79 ARG N	7.15	52.56	89.05	15.00
79 ARG CA	6.14	52.48	90.08	15.00
79 ARG CB	6.27	53.65	91.07	15.00
79 ARG CG	7.58	53.69	91.85	15.00
79 ARG CD	7.76	55.00	92.63	15.00
79 ARG NE	6.66	55.34	93.55	15.00
79 ARG CZ	6.30	54.65	94.63	15.00
79 ARG NH1	6.95	53.54	94.96	15.00
79 ARG NH2	5.21	55.01	95.31	15.00
79 ARG C	4.77	52.46	89.43	15.00
79 ARG O	3.80	52.99	90.00	15.00
BO GLY N	4.66	51.78	88.30	15.00

80 GLY CA	3.39	51.70	87.64	15.00
80 GLY C	3.39	51.50	86.16	15.00
80 GLY O	4.32	51.90	85.43	15.00
81 ILE N	2.30	50.87	85.73	15.00
81 ILE CA	2.03	50.55	84.34	15.00
81 ILE CB	2.38	49.06	83.98	15.00
81 ILE CG2	1.41	48.10	84.64	15.00
81 ILE CG1	2.48	48.88	82.46	15.00
81 ILE CD1	2.67	47.47	82.00	15.00
81 ILE C	0.57	50.92	84.01	15.00
81 ILE O	-0.33	50.73	84.82	15.00
82 ASP N	0.35	51.53	82.85	15.00
82 ASP CA	-1.01	51.93	82.43	15.00
82 ASP CB	-0.91	52.93	81.27	15.00
82 ASP CG	-0.45	54.32	81.71	15.00
82 ASP OD1	0.52	54.88	81.14	15.00
82 ASP OD2	-1.08	54.84	82.64	15.00
82 ASP C	-1.87	50.76	82.00	15.00
82 ASP O	-1.39	49.64	81.87	15.00
83 SER N	-3.16	51.03	81.87	15.00
83 SER CA	-4.11	50.02	81.42	15.00
83 SER CB	-5.54	50.23	82.00	15.00
83 SER OG	-5.97	51.58	81.87	15.00
83 SER C	-4.14	50.20	79.91	15.00
83 SER O	-3.48	51.11	79.34	15.00
84 GLU N	-4.79	49.26	79.24	15.00
84 GLU CA	-4.92	49.33	77.79	15.00
84 GLU CB	-5.77	48.16	77.25	15.00
84 GLU CG	-5.57	47.95	75.77	15.00
84 GLU CD	-4.09	47.91	75.42	15.00
84 GLU OE1	-3.52	48.88	74.89	15.00
84 GLU OE2	-3.46	46.90	75.73	15.00
84 GLU C	-5.65	50.62	77.45	15.00
84 GLU O	-5.13	51.48	76.76	15.00
85 ASP N	-6.84	50.77	78.03	15.00
85 ASP CA	-7.68	51.96	77.78	15.00
85 ASP CB	-9.03	51.85	78.51	15.00
85 ASP CG	~9.94	53.12	78.30	15.00
85 ASP OD1	-10.32	53.78	79.30	15.00
85 ASP OD2	-10.26	53.46	77.15	15.00
85 ASP C	-7.01	53.27	78.08	15.00
35 ASP O	-7.29	54.29	77.43	15.00
36 ALA N	-6.11	53.26	79.05	15.00

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86 ALA CA	-5.41	54.47	79.38	15.00
86 ALA CB	-4.88	54.38	80.80	15.00
86 ALA C	-4.27	54.73	78.39	15.00
86 ALA O	-4.00	55.89	78.07	15.00
87 TYR N	-3.69	53.66	77.83	15.00
87 TYR CA	-2.57	53.78	76.88	15.00
87 TYR CB	-1.24	53.68	77.66	15.00
87 TYR CG	0.04	54.14	77.00	15.00
87 TYR CD1	0.10	54.44	75.56	15.00
87 TYR CE1	1.32	54.83	74.98	15.00
87 TYR CD2	1.21	54.26	77.84	15.00
87 TYR CE2	2.42	54.65	77.33	15.00
87 TYR CZ	2.49	54.94	75.86	15.00
87 TYR OH	3.71	55.29	75.29	15.00
87 TYR C	-2.68	52.66	75.83	15.00
87 TYR O	-2.00	51.63	75.92	15.00
88 PRO N	-3.51	52.90	74.79	15.00
88 PRO CD	-4.53	53.97	74.75	15.00
88 PRO CA	-3.75	51.94	73.71	15.00
88 PRO CB	-4.87	52.62	72.90	15.00
88 PRO CG	-5.64	53.31	73.98	15.00
88 PRO C	-2.52	51.62	72.84	15.00
88 PRO O	-1.59	52.42	72.70	15.00
89 TYR N	-2.58	50.42	72.24	15.00
89 TYR CA	-1.54	49.87	71.38	15.00
89 TYR CB	-1.64	48.34	71.47	15.00
89 TYR CG	-0.54	47.59	70.84	15.00
89 TYR CD1	0.79	47.78	71.31	15.00
89 TYR CE1	1.83	47.13	70.72	15.00
89 TYR CD2	-0.80	46.69	69.76	15.00
89 TYR CE2	0.20	46.03	69.15	15.00
99 TYR CZ	1.53	46.24	69.61	15.00
89 TYR OH	2.55	45.63	68.93	15.00
89 TYR C	-1.73	50.30	69.92	15.00
89 TYR O	-2.81	50.16	69.38	15.00
90 VAL N	-0.65	50.73	69.28	15.00
90 VAL CA	-0.69	51.22	67.88	15.00
90 VAL CB	-0.15	52.67	67.77	15.00
90 VAL CG1	-0.81	53.58	58.80	15.00
90 VAL CG2	1.34	52.68	67.96	15.00
90 VAL C	0.11	50.36	66.90	15.00
90 VAL O	-0.09	50.40	65.68	15.00
91 GLY N	1.05	49.58	67.43	15.00

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91 GLY CA	1.85	48.75	66.55	15.00
91 GLY C	3.14		66.11	
91 GLY O	3.81	48.92		
92 GLN N	3.51		66.75	
92 GLN CA	4.75	_	66.39	
92 GLN CB	4.53		65.23	
92 GLN CG	3.31	52.97	65.48	
92 GLN CD	3.44		64.81	
92 GLN OE1	2.68	54.64	63.92	
92 GLN NE2	4.41	55.10	65.25	
92 GLN C	5.34		67.53	
92 GLN O	4.72	52.25		
93 GLU N	6.57	52.40	67.30	
93 GLU CA		53.20	68.25	
93 GLU CB	8.81	53.08	67.91	
93 GLU CG	9.33	51.65	68.15	
93 GLU CD	10.59		67.38	
93 GLU OE1	10.87	– –	66.34	15.00
93 GLU OE2	11.31		67.80	15.00
93 GLU C	6.85	54.66	68.26	15.00
93 GLU O	6.48	55.24	67.24	15.00
94 GLU N	6.79	55.20	69.46	15.00
94 GLU CA	6.36	56.56		15.00
94 GLU CB	4.83		69.62	15.00
94 GLU CG	4.07		70.04	15.00
94 GLU CD	2.76	55.78	70.70	15.00
94 GLU OE1	1.99	56.60	70.14	15.00
94 GLU OE2	2.51	55.26	71.81	15.00
94 GLU C	6.86	-	71.01	15.00
94 GLU O	7.15	56.12	71.84	15.00
95 SER N	6.98	58.29	71.23	15.00
95 SER CA	7.47	58.78		
95 SER CB		60.29		15.00
95 SER OG	6.14	60.73	71.85	15.00
95 SER C	6.78	58.06	73.70	15.00
95 SER O	5.61	<b>57</b> .59	73.59	15.00
96 CYS N	7.52	57.90	74.80	15.00
96 CYS CA	6.93	57.30	75.96	15.00
96 CYS C	5.80	58.23	76.33	15.00
96 CYS O	6.00	59.44	76.37	15.00
96 CYS CB	7.95	57.27	77.09	15.00
96 CYS SG	7.32	56.63	78.67	15.00
97 MET N	4.58	57.73	7€.42	15.00

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97 MET CA	3.46	58.59	76.84	15.00
97 MET CB	2.40	58.64	75.75	15.00
97 MET CG	2.95	58.79	74.35	15.00
97 MET SD	1.74	59.21	73.09	15.00
97 MET CE	0.27	58.48	73.83	15.00
97 MET C	2.79	58.10	78.14	15.00
97 MET 0	1.57	58.06	78.24	15.00
98 TYR N	3.59	57.79	79.15	15.00
98 TYR CA	3.09	57.31	80.45	15.00
98 TYR CB	4.25	56.91	81.37	15.00
98 TYR CG	3.71	56.31	82.64	15.00
98 TYR CD1	2.99	55.13	82.59	15.00
98 TYR CE1	2.42	54.60	83.75	15.00
98 TYR CD2	3.86	56.96	83.88	15.00
98 TYR CE2	3.30	56.45	85.03	15.00
98 TYR CZ	2.59	55.27	84.95	15.00
98 TYR OH	1.99	54.70	86.05	15.00
98 TYR C	2.17	58.29	81.19	15.00
98 TYR O	2.56	59.42	81.48	15.00
99 ASN N	0.97	57.85	81.49	15.00
99 ASN CA	-0.01	58.66	82.21	15.00
99 ASN CB	-1.39	58.64	81.52	15.00
99 ASN CG	-2.41	59.49	82.25	15.00
99 ASN OD1	-2.15	59.98	83.35	15.00
99 ASN ND2	-3.60	59.65	81.65	15.00
99 ASN C	-0.17	58.09	83.63	15.00
99 ASN O	-0.87	57.05	83.81	15.00
100 PRO N	0.31	58.84	84.66	15.00
100 PRO CD	0.81	60.23	84.63	15.00
100 PRO CA	0.20	58.38	86.04	15.00
100 PRO CB	0.80	59.53	86.83	15.00
100 PRO CG	1.64	60.28	85.79	15.00
100 PRO C	-1.25	58.14	86.50	15.00
100 PRO O	-1.49	57.37	87.41	15.00
101 THR N	-2.22	58.73	85.82	15.00
101 THR CA	-3.61	58.53	86.23	15.00
101 THR CB	-4.58	59.63	85.74	15.00
101 THR OG1	-5.12	59.28	84.45	15.00
101 THR CG2	-3.91	60.96	85.67	15.00
101 THR C	-4.09	57.21	85.69	15.00
101 THR O	-5.05	56.64	86.24	15.00
102 GLY N	~3.50	56.77	84.58	15.00
102 GLY CA	-3.90	55.51	83.98	15.00

102 GLY C	-3.31	54.30	84.66	15.00
102 GLY O	-3.63	53.16	84.25	
103 LYS N	-2.50	54.52	85.70	15.00
103 LYS CA	-1.87	53.41	86.40	15.00
103 LYS CB	-1.06	53.93		
103 LYS CG	-0.26			
103 LYS CD	-0.09			
103 LYS CE	-1.41	52.95	90.56	
103 LYS NZ	-1.62	51.48		15.00
103 LYS C	-2.90	52.38		
103 LYS O	-3.84	52.70		
104 ALA N	-2.70	51.13	86.45	15.00
104 ALA CA	-3.60	50.03	86.82	15.00
104 ALA CB	-4.23	49.45	85.59	15.00
104 ALA C	-2.81	48.95		
104 ALA O	-3.37			
105 ALA N	-1.49	49.07		15.00
105 ALA CA	-0.72	48.05	88.20	15.00
105 ALA CB	-0.65	46.80	87.34	15.00
105 ALA C	0.66	48.46	88.70	15.00
105 ALA O	1.21	49.47		15.00
106 LYS N	1.18		89.63	15.00
106 LYS CA	2.49	47.81	90.27	15.00
106 LYS CB	2.33	48.37	91.70	15.00
106 LYS CG	1.94	49.82	91.78	15.00
106 LYS CD	2.39	50.49	93.08	
106 LYS CE	3.90	50.73	93.15	15.00
106 LYS NZ	4.68	49.45	93.03	15.00
106 LYS C	3.31	46.52	90.33	15.00
106 LYS O	2.78	45.42	90.25	15.00
107 CYS N	4.58	46.66	90.67	15.00
107 CYS CA	5.44	45.50	90.73	15.00
107 CYS CB	5.96	45.28	89.31	15.00
107 CYS SG	7.11	43.91	89.12	15.00
107 CYS C	6.58	45.74	91.71	15.00
107 CYS 0	7.13	46.84	91.75	15.00
108 ARG N	6.93	44.73	92.50	15.00
108 ARG CA	8.03	44.94	93.42	15.00
108 ARG CB	7.57	44.79	94.87	15.00
108 ARG CG	7.03	43.41	95.26	15.00
108 ARG CD	6.66	43.37	96.77	15.00
108 ARG NE	6.16	42.05	97.15	15.00
108 ARG CZ	6.83	40.90	96.99	15.00

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108 ARG NH1	8.05	40.88	96.47	15.00
108 ARG NH2	6.25	39.76	97.31	
108 ARG C	9.24	44.06	93.14	15.00
108 ARG O	9.94		94.07	15.00
109 GLY N	9.47	43.78	91.85	15.00
109 GLY CA	10.58		91.49	15.00
109 GLY C	10.15	41.85	90.50	
109 GLY 0	9.05		89.95	15.00
110 TYR N	10.97	40.81	90.37	15.00
110 TYR CA	10.68	39.71	89.47	15.00
110 TYR CB	10.79	40.15	88.01	
110 TYR CG	12.20	40.50	87.61	
110 TYR CD1	12.66	41.85	87.79	
110 TYR CE1	13.95	42.21	87.41	15.00
110 TYR CD2	13.08	39.49	87.02	15.00
110 TYR CE2	14.35	39.80	86.64	15.00
110 TYR CZ	14.80	41.19	86.83	15.00
110 TYR OH	16.07		86.47	15.00
110 TYR C	11.67	38.57	89.72	15.00
110 TYR O	12.72	38.78	90.34	15.00
111 ARG N	11.36	37.37	89.26	15.00
111 ARG CA	12.26	36.23	89.43	15.00
111 ARG CB		35.41	90.67	15.00
111 ARG CG	11.78		91.94	15.00
111 ARG CD	12.04		93.15	15.00
111 ARG NE	13.43	34.92	93.18	15.00
111 ARG CZ	13.89	33.89	93.89	15.00
111 ARG NH1	15.19		93.83	15.00
111 ARG NH2	13.04		94.59	15.00
111 ARG C	12.24	35.31	88.24	15.00
111 ARG O	11.17	35.05	87.66	15.00
112 GLU N	13.43		87.90	15.00
112 GLU CA	13.63			15.00
112 GLU CB	15.09	33.97	86.40	15.00
112 GLU CG	15.46	35.25	85.64	15.00
112 GLU CD	15.38	35.07	84.12	15.00
112 GLU OE1	14.25	35.12	83.56	15.00
112 GLU OE2	16.47	34.88	83.50	15.00
112 GLU C	13.36	32.48	87.30	15.00
112 GLU O	13.00	32.27	88.44	15.00
113 ILE N	13.55	31.53	86.41	15.00
113 ILE CA	13.36	30.12	86.72	15.00
113 ILE CB	12.20	29.54	85.92	15.00

#### TABLE X 113 ILE CG2 12.22 28.02 85.90 15.00 113 ILE CG1 10.91 30.06 86.51 15.00 113 ILE CD1 9.69 29.82 85.68 15.00 113 ILE C 14.68 29.49 86.33 15.00 113 ILE 0 15.33 29.97 85.44 15.00 114 PRO N 15.20 28.55 87.13 15.00 114 PRO CD 14.60 28.07 88.38 15.00 114 PRO CA 16.49 27.87 86.88 15.00 114 PRO CB 16.38 26.62 87.77 15.00 114 PRO CG 15.73 27.17 88.97 15.00 114 PRO C 16.69 27.54 85.40 15.00 114 PRO 0 15.93 26.78 84.83 15.00 115 GLU N 17.71 28.13 84.80 15.00 115 GLU CA 18.03 28.00 83.38 15.00 115 GLU CB 19.41 28.59 83.11 15.00 115 GLU CG 19.58 29.05 81.64 15.00 115 GLU CD 20.85 29.87 81.42 15.00 115 GLU OE1 20.80 31.11 81.48 15.00 115 GLU OE2 21.91 29.28 81.19 15.00 115 GLU C 17.96 26.62 82.77 15.00 115 GLU 0 18.99 25.96 82.65 15.00 116 GLY N 16.78 26.24 82.28 15.00 116 GLY CA 16.64 24.93 81.66 15.00 116 GLY C 15.95 23.89 82.52 15.00 116 GLY O 15.82 22.73 82.13 15.00 117 ASN N 15.49 24.30 83.70 15.00 117 ASN CA 14.84 23.38 84.61 15.00 117 ASN CB 15.29 23.66 86.03 15.00 117 ASN CG 14.55 22.85 87.06 15.00 117 ASN OD1 13.47 22.36 86.81 15.00 117 ASN ND2 15.14 22.73 88.25 15.00 117 ASN C 13.34 23.50 84.47 15.00 117 ASN 0 12.71 24.37 85.05 15.00 118 GLU N 12.78 22.55 83.74 15.00 118 GLU CA 11.35 22.52 83.45 15.00 118 GLU CB 11.10 21.52 82.33 15.00 118 GLU CG 10.04 21.91 81.37 15.00 118 GLU CD 9.94 20.95 80.16 15.00 118 GLU OE1 8.82 20.50 79.88 15.00 118 GLU OE2 10.96 20.58 79.52 15.00

22.18

22.61

21.46

84.64

84.67

85.63

15.00

15.00

15.00

10.45

9.30

10.97

118 GLU C

118 GLU O

119 LYS N

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119 LYS CA	10.15		86.77	15.0
119 LYS CB	10.74	55	87.52	
119 LYS CG	9.69		88.07	
119 LYS CD	8.84		86.94	
119 LYS CE	7.78			
119 LYS NZ	7.03	17.08		
119 LYS C	10.02	22.33		
119 LYS O	9.02	22.45		-
120 ALA N	11.01	23.22		
120 ALA CA	10.99	24.43		
120 ALA CB	12.36	25.05		
120 ALA C	9.95	25.39		-
120 ALA O	9.39			15.00
121 LEU N	9.79			15.00
121 LEU CA	8.83		-	
121 LEU CB	9.06	25.87	84.28	
121 LEU CG	8.06	26.54		15.00
121 LEU CD1	8.05	28.02	83.54	15.00
121 LEU CD2	8.42		81.90	15.00
121 LEU C	7.43	25.59	86.13	15.00
121 LEU O	6.59	26.43	86.46	
122 LYS N	7.17	24.28		15.00
122 LYS CA	5.83	23.74	86.40	15.00
122 LYS CB	5.81		86.25	15.00
122 LYS CG	4.49	21.54	86.59	15.00
122 LYS CD	4.61	20.10	86.92	15.00
122 LYS CE	5.26	19.90	88.29	15.00
122 LYS NZ	5.14	18.48	88.78	15.00
122 LYS C	5.34	24.13	87.78	15.00
122 LYS 0	4.13	24.20	87.99	15.00
123 ARG N	6.27	24.36	88.71	15.00
123 ARG CA	5.90	24.76	90.06	15.00
123 ARG CB	6.95			15.00
123 ARG CG	7.05	22.94	91.45	15.00
123 ARG CD	8.15	22.86	92.43	15.00
123 ARG NE	9.44	23.25	91.82	15.00
123 ARG CZ	10.56	23.53	92.50	15.00
123 ARG NH1	10.58	23.50	93.81	15.00
123 ARG NH2	11.71	23.76	91.85	15.00
123 ARG C	5.71	26.25	90.17	15.00
123 ARG O	5.12	26.72	91.13	15.00
124 ALA N	6.31	27.00	89.25	15.00
124 ALA CA	6.16	28.42	89.30	15.00

124 ALA CB	7.22	29.12	88.43	15.00
124 ALA C	4.78	28.73	88.81	
124 ALA O	4.06			
125 VAL N	4.37			
125 VAL CA	3.06		87.21	15.00
125 VAL CB	2.82	27.94	85.72	
125 VAL CG1	4.09	27.54		
125 VAL CG2	1.70			
125 VAL C	2.03		88.15	15.00
125 VAL 0	0.89	28.28	88.17	15.00
126 ALA N	2.42	26.86	88.98	15.00
126 ALA CA	1.47	26.25		
126 ALA CB	1.94			15.00
126 ALA C	1.32	27.11	91.12	15.00
126 ALA O	0.22	27.27	91.63	15.00
127 ARG N	2.42	27.71	91.55	15.00
127 ARG CA	2.42	28.50	92.77	
127 ARG CB	3.67	28.22	93.60	15.00
127 ARG CG	3.74	26.79	94.07	15.00
127 ARG CD	5.07	26.45	94.67	15.00
127 ARG NE	5.02	25.01	94.96	15.00
127 ARG CZ	6.03	24.24	95.34	15.00
127 ARG NH1	5.84	22.94	95.55	15.00
127 ARG NH2	7.24	24.73	95.54	15.00
127 ARG C	2.30	29.98	92.61	15.00
127 ARG 0	2.15	30.68	93.60	15.00
128 VAL N	2.46	30.47	91.38	15.00
128 VAL CA	2.37	31.91	91.11	15.00
128 VAL CB	3.70	32.49	90.59	15.00
128 VAL CG1	3.62	33.98	90.47	15.00
128 VAL CG2	4.87	32.10	91.53	15.00
128 VAL C	1.29	32.25		
128 VAL 0	0.47	33.10		15.00
129 GLY N	1.30	31.60	88.94	15.00
129 GLY CA	0.31	31.85	87.91	15.00
129 GLY C	1.10	32.00	86.61	15.00
129 GLY 0	2.27	31.60	86.57	15.00
130 PRO N	0.52	32.56	85.53	15.00
130 PRO CD	-0.89	32.96	85.30	15.00
130 PRO CA	1.27	32.70	84.30	15.00
130 PRO CB	0.38	33.61		15.00
130 PRO CG			83.81	
130 PRO C	2.68	33.24	84.44	15.00

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130 PRO O	2.93	34.17	85.20	15.00
131 VAL N	3.59	32.56	83.75	15.00
131 VAL CA	5.01	32.88	83.76	15.00
131 VAL CB	5.79		84.27	15.00
131 VAL CG1	7.25	31.88	84.32	15.00
131 VAL CG2	5.29	31.20	85.62	15.00
131 VAL C	5.46	33.28	82.34	
131 VAL 0	4.93	32.82		15.00
132 SER N	6.29	34.30	82.27	15.00
132 SER CA	6.77	34.75	80.98	15.00
132 SER CB	7.15	36.24	81.04	15.00
132 SER OG	5.98	36.98	81.39	15.00
132 SER C	7.92	33.89		15.00
132 SER 0	9.01	33.83	81.08	15.00
133 VAL N	7.65	33.20	79.38	15.00
133 VAL CA	8.65	32.34	78.76	15.00
133 VAL CB	8.09	30.90		15.00
133 VAL CG1	7.58	30.35		15.00
133 VAL CG2	6.97	30.91	77.61	15.00
133 VAL C	9.05	32.84	77.37	15.00
133 VAL 0	8.48	33.81	76.85	15.00
134 ALA N	10.08	32.21	76.81	15.00
134 ALA CA	10.60	32.51	75.49	15.00
134 ALA CB	11.89	33.29	75.59	15.00
134 ALA C	10.85	31.16	74.84	15.00
134 ALA O	11.46	30.30	75.47	15.00
135 ILE N	10.35	30.97	73.62	15.00
135 ILE CA	10.52	29.73	72.86	15.00
135 ILE CB	9.18	28.95	72.80	15.00
135 ILE CG2	8.71	28.60	74.21	15.00
135 ILE CG1	8.13	29.80	72.09	15.00
135 ILE CD1	6.78	29.14	72.00	15.00
135 ILE C	11.04	30.04	71.44	15.00
135 ILE O	11.30	31.20	71.08	15.00
136 ASP N	11.28	28.98	70.67	15.00
136 ASP CA	11.71	29.13	69.30	15.00
136 ASP CB	12.68	28.01	68.94	15.00
136 ASP CG	13.21	28.11	67.50	15.00
136 ASP OD1	13.77	27.12	67.00	15.00
136 ASP OD2	13.08	29.18	66.88	15.00
136 ASP C	10.45	28.94	68.51	15.00
136 ASP 0	9.90	27.82	68.48	15.00
137 ALA N	9.98	30.01	67.87	15.00

137 ALA CA	8.78	29.93	67.06	15.00
137 ALA CB	7.74	30.91	67.56	15.00
137 ALA C	9.09	30.18	65.58	15.00
137 ALA O	8.27	30.79	64.88	15.00
138 SER N	10.22	29.67	65.10	
138 SER CA	10.66	29.83	63.72	
138 SER CB	12.18	29.81	63.63	15.00
138 SER OG	12.67	28.54	64.01	15.00
138 SER C	10.15	28.77	62.77	15.00
138 SER O	9.81	29.07	61.62	15.00
139 LEU N	10.12	27.53	63.26	15.00
139 LEU CA	9.66	26.42	62.45	15.00
139 LEU CB	9.74	25.13	63.29	15.00
139 LEU CG	11.05	24.33	63.34	15.00
139 LEU CD1	12.24	25.24	63.67	
139 LEU CD2	10.92	23.21	64.38	15.00
139 LEU C	8.24	26.59	61.88	15.00
139 LEU O	7.32	26.95	62.60	15.00
140 THR N	8.08	26.30	60.58	15.00
140 THR CA	6.80	26.39	59.88	15.00
140 THR CB	6.88	25.76	58.50	15.00
140 THR OG1	8.18	25.97	57.94	15.00
140 THR CG2	5.86	26.37	57.60	15.00
140 THR C	5.76	25.62	60.67	15.00
140 THR O	4.67	26.12	60.88	15.00
141 SER N	6.13	24.45	61.17	15.00
141 SER CA	5.20	23.63	61.94	15.00
141 SER CB	5.79	22.28	62.28	15.00
141 SER OG	7.00	22.40	62.97	15.00
141 SER C	4.65	24.28	63.18	15.00
141 SER O	3.60	23.89	63.66	15.00
142 PHE N	5.35	25.26	63.72	15.00
142 PHE CA	4.82	25.94	64.90	15.00
142 PHE CB	5.94	26.71	65.64	15.00
142 PHE CG	5.46	27.49	66.86	15.00
142 PHE CD1	5.60	26.99	68.12	15.00
142 PHE CD2	4.89	28.74	66.71	15.00
142 PHE CE1	5.15	27.76	69.25	15.00
142 PHE CE2	4.46	29.49	67.82	15.00
142 PHE CZ	4.59	29.00	69.07	15.00
142 PHE C	3.74	26.88	64.42	15.00
142 PHE 0	2.63	26.88	64.93	15.00
143 GLN N	4.08	27.58	63.36	15.00

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143 GLN CA	3.24	28.59	62.73	15.00
143 GLN CB	4.04	29.36	61.69	
143 GLN CG	5.38	29.87	62.23	
143 GLN CD	6.19	30.64	61.16	
143 GLN OE1	5.83	31.75	60.75	
143 GLN NE2	7.25	30.01	60.64	
143 GLN C	1.95	28.10	62.12	15.00
143 GLN O	0.99	28.86	62.08	
144 PHE N	1.91	26.90	61.55	
144 PHE CA	0.61	26.44		15.00
144 PHE CB	0.73	25.77	59.63	15.00
144 PHE CG	1.72	24.62	59.58	15.00
144 PHE CD1	2.76	24.61	58.69	15.00
144 PHE CD2	1.58	23.50	60.36	
144 PHE CE1	3.60	23.51	58.60	
144 PHE CE2	2.47	22.41	60.22	15.00
144 PHE CZ	3.44	22.44	59.35	15.00
144 PHE C	-0.14	25.50	61.98	15.00
144 PHE O	-1.10	24.82	61.59	15.00
145 TYR N	0.31	25.47	63.24	15.00
145 TYR CA	-0.31	24.65	64.26	15.00
145 TYR CB	0.32	24.91	65.65	15.00
145 TYR CG	-0.47	24.34	66.81	15.00
145 TYR CD1	-0.26	23.01	67.22	15.00
145 TYR CE1	-0.98	22.48	68.28	15.00
145 TYR CD2	-1.42	25.12	67.48	15.00
145 TYR CE2	-2.15	24.61	68.54	15.00
145 TYR CZ	-1.93	23.28	68.94	15.00
145 TYR OH	-2.67	22.83	70.02	15.00
145 TYR C	-1.80	24.98	64.30	15.00
145 TYR O	-2.24		64.03	15.00
146 SER N	-2.60		64.69	15.00
146 SER CA	-4.06	24.19	64.77	15.00
146 SER CB	-4.67	23.67	63.48	15.00
146 SER OG	-4.29	22.33	63.27	15.00
146 SER C	-4.67	23.42	65.95	15.00
146 SER O	-5.59	23.89	66.62	15.00
147 LYS N	-4.20	22.20	66.13	15.00
147 LYS CA	-4.67	21.36	67.20	15.00
147 LYS CB	-5.96	20.64	66.83	15.00
147 LYS CG	-5.81	19.47	65.90	15.00
147 LYS CD	-7.06	18.58	65.92	15.00
147 LYS CE	-6.91	17.41	64.96	15.00

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147 LYS NZ	-8.13		64.99	15.00
147 LYS C	-3.60	20.36	67.62	15.00
147 LYS 0	-2.62	20.16	66.90	15.00
148 GLY N	-3.82	19.73	68.78	15.00
148 GLY CA	-2.88	18.75	69.30	15.00
148 GLY C	-1.76	19.26	70.20	15.00
148 GLY O	-1.75	20.39	70.59	15.00
149 VAL N	-0.79	18.40	70.52	
149 VAL CA	0.33	18.81	71.35	15.00
149 VAL CB	0.58	17.89	72.57	15.00
149 VAL CG1	1.74	18.42	73.41	15.00
149 VAL CG2	-0.68	17.75	73.42	15.00
149 VAL C	1.59	18.86	70.48	
149 VAL O	2.06	17.84	69.96	
150 TYR N	2.13	20.07	70.35	15.00
150 TYR CA	3.32	20.33	69.53	15.00
150 TYR CB	3.50	21.83	69.29	15.00
150 TYR CG	4.59	22.21	68.31	15.00
150 TYR CD1	4.47	22.01	66.95	15.00
150 TYR CE1	5.48	22.37	66.04	15.00
150 TYR CD2	5.74	22.78	68.72	15.00
150 TYR CE2	6.75	23.13	67.81	15.00
150 TYR CZ	6.61	22.93	66.48	15.00
150 TYR OH	7.57	23.30	65.60	15.00
150 TYR C	4.60	19.72	70.12	15.00
150 TYR O	4.76	19.60	71.37	15.00
151 TYR N	5.44	19.24	69.21	15.00
151 TYR CA	6.74	18.68	69.55	15.00
151 TYR CB	6.62	17.30	70.22	15.00
151 TYR CG	7.96	16.61	70.42	15.00
151 TYR CD1	9.15	17.34	70.60	15.00
151 TYR CEL	10.41	16.67	70.84	15.00
151 TYR CD2	8.02	15.21	70.47	15.00
151 TYR CE2	9.24	14.52	70.71	15.00
151 TYR CZ	10.43	15.27	70.89	15.00
151 TYR OH	11.60	14.61	71.14	15.00
151 TYR C	7.38	18.53	68.19	15.00
151 TYR O	6.82	17.86	67.31	15.00
152 ASP N	8.47	19.2€	67.98	15.00
152 ASP CA	9.18	19.22	66.71	15.00
152 ASP CB	883	20.40	65.81	15.00
152 ASP CG	9.30	20.15	64.37	15.00
152 ASP CD1	10.54	20.06	64.18	15.00

152 ASP C 152 ASP O 11.34 20.21 67.14 15.0 153 GLU N 11.24 17.97 66.94 15.0 153 GLU CA 12.64 17.77 67.18 15.0 153 GLU CB 13.00 16.30 66.88 15.0 153 GLU CB 13.00 16.30 66.88 15.0 153 GLU CB 11.10 15.02 65.83 15.0 153 GLU OE1 11.10 15.02 65.83 15.0 153 GLU OE1 11.10 13.96 66.53 15.0 153 GLU OE2 10.06 15.52 65.30 15.0 153 GLU OE 13.60 18.73 66.48 15.0 153 GLU O 14.75 18.90 66.91 15.0 154 SER N 13.17 19.30 65.35 15.0 154 SER CA 14.02 20.24 64.60 15.0 154 SER CB 13.49 20.49 63.17 15.0 154 SER CB 13.49 20.49 63.17 15.0 154 SER C 14.13 21.57 65.34 15.0 155 CYS N 13.37 21.72 66.43 15.0 155 CYS C 14.64 23.18 67.97 15.0 155 CYS C 14.64 68.16 15.0 156 ASN C 16.31 24.44 68.16 15.0 156 ASN C 16.31 24.44 68.16 15.0 156 ASN C 16.31 24.74 68.81 15.0 156 ASN C 16.32 25.46 67.80 15.0 156 ASN C 16.22 25.55 70.11 15.0 156 ASN C 16.22 25.55 70.11 15.0 157 SER N 16.27 24.83 71.22 15.0 157 SER C 17.20 25.46 67.80 15.0 157 SER C 17.22 26.80 70.09 15.0 157 SER C 17.22 26.80 70.99 15.0 158 ASP C 16.99 27.51 73.49 15.0 158 ASP C 16.99 27.51 73.49 15.0 158 ASP C 16.99 27.51 73.49 15.0 158 ASP CB 20.71 26.91 71.57 15.00					T. WLL /	•	
152 ASP 0	152	2 ASI	OD2	8.42	20.06	63.48	15.00
153 GLU N 11.24 17.97 66.94 15.00 153 GLU CA 12.64 17.77 67.18 15.00 153 GLU CB 13.00 16.30 66.88 15.0 153 GLU CG 12.45 15.71 65.61 15.0 153 GLU CD 11.10 15.02 65.83 15.00 153 GLU OE1 11.10 13.96 66.53 15.00 153 GLU OE2 10.06 15.52 65.30 15.00 153 GLU C 13.60 18.73 66.48 15.00 153 GLU O 14.75 18.90 66.91 15.00 154 SER N 13.17 19.30 65.35 15.00 154 SER CA 14.02 20.24 64.60 15.00 154 SER CB 13.49 20.49 63.17 15.00 154 SER C 13.06 19.29 62.54 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS CB 12.15 22.99 68.17 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CB 17.20 25.46 67.80 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER C 16.22 25.55 70.11 15.00 157 SER C 16.22 25.44 72.55 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP C 16.99 27.51 73.49 15.00 158 ASP C 16.99 27.51 73.49 15.00 158 ASP C 20.91 25.47 72.03 15.00			C	10.69	19.18	66.97	15.00
153 GLU CA			0	11.34	20.21	67.14	15.00
153 GLU CB			J N	11.24	17.97	66.94	15.00
153 GLU CG	153	GLU	CA	12.64	17.77	67.18	15.00
153 GLU CG	153	GLU	CB	13.00	16.30	66.88	15.00
153 GLU CD	153	GLU	CG	12.45	15.71	65.61	15.00
153 GLU OE1	153	GLU	CD	11.10	15.02	65.83	15.00
153 GLU C	153	GLU	OE1	11.10	13.96	66.53	15.00
153 GLU C 153 GLU O 14.75 18.90 66.91 15.00 154 SER N 13.17 19.30 65.35 15.00 154 SER CA 14.02 20.24 64.60 15.00 154 SER CB 13.49 20.49 63.17 15.00 154 SER CB 13.06 19.29 62.54 15.00 154 SER C 14.13 21.57 65.34 15.00 154 SER C 14.13 21.57 65.34 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS CA 13.33 22.98 67.21 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS SG 11.63 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CB 16.51 24.37 73.60 15.00 158 ASP CB 20.71 26.91 71.57 15.00	153	GLU	OE2	10.06	15.52	65.30	15.00
153 GLU O 14.75 18.90 66.91 15.00 154 SER N 13.17 19.30 65.35 15.00 154 SER CA 14.02 20.24 64.60 15.00 154 SER CB 13.49 20.49 63.17 15.00 154 SER OG 13.06 19.29 62.54 15.00 154 SER C 14.13 21.57 65.34 15.00 154 SER O 14.95 22.40 65.00 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS CA 13.33 22.98 67.21 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS CB 12.15 22.99 68.17 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CA 16.31 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CD 18.80 25.83 69.59 15.00 156 ASN CD 18.80 25.83 69.59 15.00 156 ASN O 16.12 26.80 70.09 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.55 70.11 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00	153	GLU	C	13.60	18.73	66.48	15.00
154 SER CA 14.02 20.24 64.60 15.00 154 SER CB 13.49 20.49 63.17 15.00 154 SER OG 13.06 19.29 62.54 15.00 154 SER C 14.13 21.57 65.34 15.00 154 SER O 14.95 22.40 65.00 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS CA 13.33 22.98 67.21 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS CB 12.15 22.99 68.17 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CA 16.31 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00	153	GLU	0	14.75	18.90	66.91	15.00
154 SER CB	154	SER	N	13.17	19.30	65.35	15.00
154 SER OG	154	SER	CA	14.02	20.24	64.60	15.00
154 SER C 14.13 21.57 65.34 15.00 154 SER O 14.95 22.40 65.00 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS CA 13.33 22.98 67.21 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS SG 11.63 24.66 68.64 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CA 16.31 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 156 ASN C 16.22 25.55 70.11 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00	154	SER	CB	13.49	20.49	63.17	15.00
154 SER C 14.13 21.57 65.34 15.00 154 SER O 14.95 22.40 65.00 15.00 155 CYS N 13.37 21.72 66.43 15.00 155 CYS CA 13.33 22.98 67.21 15.00 155 CYS C 14.64 23.18 67.97 15.00 155 CYS O 15.25 22.21 68.44 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS SG 11.63 24.66 68.64 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN ND2 19.18 26.80 67.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 156 ASN C 16.22 25.55 70.11 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00	154	SER	OG	13.06	19.29	62.54	15.00
155 CYS N	154	SER	C	14.13	21.57	65.34	15.00
155 CYS CA	154	SER	0	14.95	22.40	65.00	15.00
155 CYS C 14.64 23.18 67.97 15.00 155 CYS O 15.25 22.21 68.44 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS SG 11.63 24.66 68.64 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CA 16.31 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN ND2 19.18 26.80 67.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 156 ASN O 16.12 26.80 70.09 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00	155	CYS	N	13.37	21.72	66.43	15.00
155 CYS O 15.25 22.21 68.44 15.00 155 CYS CB 12.15 22.99 68.17 15.00 155 CYS SG 11.63 24.66 68.64 15.00 156 ASN N 15.04 24.44 68.16 15.00 156 ASN CA 16.31 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CG 18.46 26.05 68.41 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN ND2 19.18 26.80 67.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 156 ASN C 16.22 25.55 70.11 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00	155	CYS	CA	13.33	22.98	67.21	15.00
155 CYS CB	155	CYS	С	14.64	23.18	67.97	15.00
155 CYS SG	155	CYS	0	15.25	22.21	68.44	15.00
156 ASN N	155	CYS	CB	12.15	22.99	68.17	15.00
156 ASN CA 16.31 24.74 68.81 15.00 156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CG 18.46 26.05 68.41 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN ND2 19.18 26.80 67.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 156 ASN C 16.22 25.55 70.11 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.91 25.47 72.03 15.00	155	CYS	SG	11.63	24.66	68.64	15.00
156 ASN CB 17.20 25.46 67.80 15.00 156 ASN CG 18.46 26.05 68.41 15.00 156 ASN OD1 18.80 25.83 69.59 15.00 156 ASN ND2 19.18 26.80 67.59 15.00 156 ASN C 16.22 25.55 70.11 15.00 156 ASN O 16.12 26.80 70.09 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CB 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER C 17.22 26.59 72.72 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00	156	ASN	N	15.04	24.44	68.16	15.00
156 ASN CG	156	ASN	CA	16.31	24.74	68.81	15.00
156 ASN OD1	156	ASN	CB	17.20	25.46	67.80	15.00
156 ASN ND2	156	ASN	CG	18.46	26.05	68.41	15.00
156 ASN C 16.22 25.55 70.11 15.00 156 ASN O 16.12 26.80 70.09 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CG 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CB 20.71 26.91 71.57 15.00	156	asn	OD1	18.80	25.83	69.59	15.00
156 ASN O 16.12 26.80 70.09 15.00 157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CG 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	156	ASN	ND2	19.18	26.80	67.59	15.00
157 SER N 16.27 24.83 71.22 15.00 157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER CG 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	156	asn	C	16.22	25.55	70.11	15.00
157 SER CA 16.22 25.44 72.55 15.00 157 SER CB 16.51 24.37 73.60 15.00 157 SER OG 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	156	asn	0	16.12	26.80	70.09	15.00
157 SER CB 16.51 24.37 73.60 15.00 157 SER OG 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	157	SER	N	16.27	24.83	71.22	15.00
157 SER OG 15.49 23.40 73.56 15.00 157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	157	SER	CA	16.22	25.44	72.55	15.00
157 SER C 17.22 26.59 72.72 15.00 157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	157	SER	CB	16.51	24.37	73.60	15.00
157 SER O 16.99 27.51 73.49 15.00 158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	157	SER	OG	15.49	23.40	73.56	15.00
158 ASP N 18.35 26.52 72.02 15.00 158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	157	SER	C	17.22	26.59	72.72	15.00
158 ASP CA 19.40 27.52 72.06 15.00 158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	157	SER	0	16.99	27.51	73.49	15.00
158 ASP CB 20.71 26.91 71.57 15.00 158 ASP CG 20.91 25.47 72.03 15.00	158	ASP	N	18.35	26.52	72.02	15.00
158 ASP CG 20.91 25.47 72.03 15.00	158	ASP	CA	19.40	27.52	72.06	15.00
	158	ASP	CB	20.71	26.91	71.57	15.00
158 ASP OD1 21.16 24.60 71.16 15.00	158	ASP	CG	20.91	25.47		15.00
	158	ASP	OD1	21.16	24.60	71.16	15.00

		IABLE	<b>^</b>	
158 ASP OD2	20.81	25.18	73.26	15.00
158 ASP C	19.05	28.77	71.26	15.00
158 ASP O	19.69	29.81	71.37	15.00
159 ASN N	18.04	28.67	70.40	15.00
159 ASN CA	17.64	29.82	69.62	15.00
159 ASN CB	17.77	29.53	68.12	15.00
159 ASN CG	17.54	30.78	67.23	15.00
159 ASN OD1	17.33	30.67	66.00	15.00
159 ASN ND2	17.63	31.96	67.83	15.00
159 ASN C	16.22	30.27	69.99	15.00
159 ASN 0	15.23	29.87	69.36	15.00
160 LEU N	16.12	31.10	71.03	15.00
160 LEU CA	14.84	31.64	71.49	15.00
160 LEU CB	14.88	31.87	73.00	15.00
160 LEU CG	15.40	30.70	73.88	15.00
160 LEU CD1	15.23	31.09	75.33	15.00
160 LEU CD2	14.68	29.38	73.59	15.00
160 LEU C	14.66	32.96	70.75	15.00
160 LEU O	15.56	33.76	70.75	15.00
161 ASN N	13.52	33.17	70.11	15.00
161 ASN CA	13.28	34.40	69.36	15.00
161 ASN CB	13.53	34.18	67.85	15.00
161 ASN CG	12.91	32.90	67.32	15.00
161 ASN OD1	11.68	32.70	67.33	15.00
161 ASN ND2	13.78	32.01	66.83	15.00
161 ASN C	11.86	34.98	69.54	15.00
161 ASN 0	11.57	36.09	69.09	15.00
162 HIS N	10.99	34.28	70.26	15.00
162 HIS CA	9.66	34.79	70.41	15.00
162 HIS CB	8.74	34.04	69.45	15.00
162 HIS CG	7.37	34.62	69.35	15.00
162 HIS CD2	6.94	35.88	69.10	15.00
162 HIS ND1	6.24	33.84	69.45	15.00
162 HIS CE1	5.17	34.59	69.25	15.00
162 HIS NE2	5.57	35.83	69.03	15.00
162 HIS C	9.28	34.53	71.85	15.00
162 HIS O	9.61	33.48	72.39	15.00
163 ALA N	8.70	35.56	72.47	15.00
163 ALA CA	8.26	35.51	73.85	15.00
163 ALA CB	8.50	36.80	74.53	15.00
163 ALA C	6.78	35.24	73.87	15.00
163 ALA O	6.02	35.80	73.09	15.00
164 VAL N	6.39	34.38	74.78	15.00

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164 VAL CA	5.01		74.97	15.00
164 VAL CB	4.69		74.29	
164 VAL CG1	4.67		72.75	
164 VAL CG2	5.73	31.64	74.69	
164 VAL C	4.72	33.94	76.48	15.00
164 VAL O	5.55		77.30	15.00
165 LEU N	3.60	33.31		
165 LEU CA	3.27	33.21		
165 LEU CB	2.31	34.36		15.00
165 LEU CG	1.52	34.51		15.00
165 LEU CD1	2.25	35.21		15.00
165 LEU CD2	0.26	35.25	79.65	
165 LEU C	2.65	31.88	78.63	
165 LEU O	1.68	31.50		
166 ALA N	3.25	31.15		
166 ALA CA	2.73	29.85		15.00
166 ALA CB	3.79	29.05	80.72	15.00
166 ALA C	1.57	30.12	80.98	15.00
166 ALA O	1.76	30.63	82.05	
167 VAL N	0.38	29.73	80.56	
167 VAL CA	-0.88	_		15.00
167 VAL CB	-1.94	30.48	80.19	15.00
167 VAL CG1	-3.36	30.02	80.45	15.00
167 VAL CG2	-1.88	32.03	80.10	15.00
167 VAL C	-1.36	28.70	82.03	15.00
167 VAL O	-2.35	28.75		15.00
168 GLY N	-0.67	27.57	81.88	15.00
168 GLY CA	-1.07	26.35	82.56	15.00
168 GLY C	-0.36	25.13	82.00	15.00
168 GLY O	0.52	25.29	81.17	15.00
169 TYR N	-0.68	23.94	82.49	15.00
169 TYR CA	-0.03		81.98	15.00
169 TYR CB	1.33	22.45	82.69	15.00
169 TYR CG	1.24	22.33	84.21	15.00
169 TYR CD1	0.66	21.21	84.82	15.00
169 TYR CE1	0.51	21.16	86.19	15.00
169 TYR CD2	1.67	23.36	85.02	15.00
169 TYR CE2	1.53	23.30	86.36	15.00
169 TYR CZ	0.94	22.20	86.95	15.00
169 TYR OH	0.76	22.18	88.32	15.00
169 TYR C	-0.97	21.62	82.29	15.00
169 TYR O	-1.89	21.79	83.03	15.00
170 GLY N	-0.71	20.47	81.70	15.00

170 GLY CA	-1.54		81.93	15.00
170 GLY C	-1.19	18.20	80.97	15.00
170 GLY O	-0.05	18.05	80.56	15.00
171 ILE N	-2.19	17.42	80.63	15.00
171 ILE CA	-1.99	16.31	79.71	15.00
171 ILE CB	-1.71	14.98	80.49	15.00
171 ILE CG2	-2.84	14.65	81.48	15.00
171 ILE CG1	-1.47	13.80	79.57	15.00
171 ILE CD1	-1.37	12.49	80.30	15.00
171 ILE C	-3.23	16.19	78.81	15.00
171 ILE O	-4.29	16.77	79.10	15.00
172 GLN N	-3.09	15.50	77.69	15.00
172 GLN CA	-4.21	15.35	76.77	15.00
172 GLN CB	-3.96	16.16	75.48	15.00
172 GLN CG	<b>-</b> 5.15	16.20	74.49	15.00
172 GLN CD	-5.05	17.33	73.50	15.00
172 GLN OE1	-4.76	18.47	73.87	15.00
172 GLN NE2	-5.27	17.03	72.22	15.00
172 GLN C	-4.51	13.86	76.49	15.00
172 GLN 0	-5.23	13.22	77.27	15.00
173 LYS N	-4.00	13.30	75.41	15.00
173 LYS CA	-4.27	11.89	75.22	15.00
173 LYS CB	-4.76	11.64	73.81	15.00
173 LYS CG	-6.07	12.36	73.52	15.00
173 LYS CD	-6.45	12.30	72.03	15.00
173 LYS CE	-5.44	13.01	71.15	15.00
173 LYS NZ	-5.85	12.96	69.72	15.00
173 LYS C	-2.97	11.18	75.51	15.00
173 LYS O	-2.29	10.72	74.59	15.00
174 GLY N	-2.59	11.22	76.78	15.00
174 GLY CA	-1.36	10.60	77.21	15.00
174 GLY C	-0.18	11.54	77.14	15.00
174 GLY O	0.88	11.27	77.72	15.00
175 ASN N	-0.34	12.66	76.42	15.00
175 ASN CA	0.75	13.62	76.27	15.00
175 ASN CB	0.84	14.07	74.82	15.00
175 ASN CG	1.30	12.97	73.90	15.00
175 ASN OD1	2.46	12.52	73.96	15.00
175 ASN ND2	0.38	1.2.50	73.05	15.00
175 ASN C	0.75	14.84	77.17	15.00
175 ASN 0	-0.24	15.61	77.23	15.00
176 LYS N	1.91	15.07	77.78	15.00
176 LYS CA	2.12	16.20	78.66	15.00

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176 LYS CB	3.36		79.53	15.00
176 LYS CG	3.19			
176 LYS CD	4.24	14.85		
176 LYS CE	4.17			
176 LYS NZ	2.95			
176 LYS C	2.25	17.46		
176 LYS 0	2.71			
177 HIS N	1.74		78.38	
177 HIS CA	1.85	19.81	77.66	
177 HIS CB	0.73	19.92	76.63	
177 HIS CG	-0.62	20.17		15.00
177 HIS CD2	-1.20	21.32	77.65	15.00
177 HIS ND1	-1.56	19.18	77.38	15.00
177 HIS CE1	-2.67	19.70	77.89	
177 HIS NE2	-2.47	21.00	78.06	
177 HIS C	1.83	21.03	78.53	15.00
177 HIS O	1.38	20.97	79.68	15.00
178 TRP N	2.20	22.13	77.89	15.00
178 TRP CA	2.21	23.47	78.45	15.00
178 TRP CB	3.57	24.15	78.26	15.00
178 TRP CG	4.71	23.55	78.98	15.00
178 TRP CD2	4.98	23.62	80.41	15.00
178 TRP CE2	6.14	22.88	80.64	15.00
178 TRP CE3	4.33	24.22	81.52	15.00
178 TRP CD1	5.70	22.82	78.45	15.00
178 TRP NE1	6.56	22.40	79.44	15.00
178 TRP CZ2	6.67	22.72	81.91	15.00
178 TRP CZ3	4.86	24.05	82.77	15.00
178 TRP CH2	6.01		82.96	15.00
178 TRP C 178 TRP O	1.18	24.24	77.64	15.00
-	1.14	24.11	76.42	15.00
	0.33	25.01	78.30	15.00
	-0.64	25.81	77.55	15.00
	-1.90	26.08	78.34	15.00
179 ILE CG2	-2.77	27.07	77.61	15.00
179 ILE CG1	-2.54	24.73	78.71	15.00
179 ILE CD1	-3.79	24.79	79.53	15.00
179 ILE C	0.03	27.14	77.31	15.00
179 ILE O	0.36	27.82	78.27	15.00
180 ILE N	0.23	27.49	76.05	15.00
180 ILE CA	0.89	28.72	75.68	15.00
180 ILE CB	2.09	28.49	74.78	15.00
180 ILE CG2	2.73	29.81	74.51	15.00

180 ILE CG1	2 00	22		
180 ILE CD1	3.09		75.46	
180 ILE C	3.92		76.47	
180 ILE O	-0.04		75.03	15.00
	-0.88	_	74.22	15.00
181 LYS N	0.06	30.94	75.44	15.00
181 LYS CA	-0.80	31.95	74.91	
181 LYS CB	-1.26	32.84		15.00
181 LYS CG	-1.94			15.00
181 LYS CD	-2.47			15.00
181 LYS CE	-2.92	36.22	76.11	15.00
181 LYS NZ	-3.47	37.09	77.17	15.00
181 LYS C	0.15	32.72	74.03	15.00
181 LYS O	1.19	33.15	74.52	15.00
182 ASN N	-0.15	32.81	72.75	15.00
182 ASN CA	0.73	33.49	71.81	15.00
182 ASN CB	0.96	32.61	70.59	15.00
182 ASN CG	1.97	33.17	69.65	15.00
182 ASN OD1	2.31	34.34	69.73	15.00
182 ASN ND2	2.42	32.35	68.71	15.00
182 ASN C	0.01	34.76	71.38	15.00
182 ASN 0	-1.21	34.83	71.39	15.00
183 SER N	0.75	35.80	71.01	15.00
183 SER CA	0.09	37.03	70.61	
183 SER CB	0.81	38.23	71.20	15.00
183 SER OG	2.19	38.09	71.04	15.00
183 SER C	-0.09	37.18	69.10	15.00
183 SER O	0.19	38.25	68.56	15.00
184 TRP N	-0.58	36.16	68.41	15.00
184 TRP CA	-0.73	36.29	66.97	15.00
184 TRP CB	0.00	35.18	66.21	15.00
184 TRP CG	1.47	35.38	66.20	15.00
184 TRP CD2	2.44	34.39	65.92	15.00
184 TRP CE2	3.69	35.01	65.97	15.00
184 TRP CE3	2.38	33.02	65.61	15.00
184 TRP CD1	2.13	36.54	66.41	15.00
184 TRP NE1	3.47	36.33	66.27	15.00
184 TRP CZ2	4.88	34.32	65.74	15.00
184 TRP CZ3	3.56	32.33	65.37	15.00
184 TRP CH2	4.79	32.98	65.44	15.00
184 TRP C	-2.17	36.42	66.57	15.00
184 TRP O	-2.53	36.20	65.42	15.00
185 GLY N	-2.97	36.92	67.51	15.00
185 GLY CA	-4.37	37.14	67.23	15.00
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185 GLY C	-5.06	35.83	67.50	15.00
185 GLY O	-4.46	34.76	67.64	15.00
186 GLU N	-6.37	35.94	67.57	15.00
186 GLU CA	-7.23	34.80	67.84	15.00
186 GLU CB	-8.63	35.32	68.26	15.00
186 GLU CG	-9.57	34.27	68.68	15.00
186 GLU CD	-10.89	34.86	69.09	15.00
186 GLU OE1	-11.68	35.25	68.20	15.00
186 GLU OE2	-11.14	34.91	70.31	15.00
186 GLU C	-7.36	33.76	66.75	15.00
186 GLU O	-7.74	32.62	67.03	15.00
187 ASN N	-7.00	34.14	65.52	15.00
187 ASN CA	-7.11	33.25	64.36	15.00
187 ASN CB	-7.33	34.07	63.08	15.00
187 ASN CG	-8.45	33.50	62.21	15.00
187 ASN OD1	-8.22	33.17		15.00
187 ASN ND2	-9.66	33.42	62.75	15.00
187 ASN C	-5.92	32.36	64.14	15.00
187 ASN O	-5.90	31.63	63.13	15.00
188 TRP N	-4.89	32.50	64.96	15.00
188 TRP CA	-3.73	31.66	64.83	15.00
188 TRP CB	-2.53	32.41	65.32	15.00
188 TRP CG	-1.34	31.51	65.25	15.00
188 TRP CD2	-0.66	30.87	66.35	15.00
188 TRP CE2	0.34	30.12	65.83	15.00
188 TRP CE3	-0.81	30.87	67.72	15.00
188 TRP CD1	-0.72	31.11	64.13	15.00
188 TRP NE1	0.30	30.26	64.46	15.00
188 TRP CZ2	1.18	29.39	66.64	15.00
188 TRP CZ3	0.04	30.13	68.52	15.00
188 TRP CH2	0.98	29.43	68.01	15.00
188 TRP C	-3.93	30.46	65.72	15.00
188 TRP 0	-4.69	30.59	66.69	15.00
189 GLY N	-3.29	29.33	65.40	15.00
189 GLY CA	-3.42	28.14	66.22	15.00
189 GLY C	-4.80	27.88	66.77	15.00
189 GLY O	-5.82	28.15	66.15	15.00
190 ASN N	-4.87	27.38	67.98	15.00
190 ASN CA	-6.16	27.08	68.58	15.00
190 ASN CB	-6.04	25.89	69.53	15.00
190 ASN CG	-7.37	25.28	69.86	
190 ASN OD1	-8.42	25.95		15.00
190 ASN ND2	-7.34	24.00	70.23	15.00

190 ASN C	-6.70	28.34	69.26	15.00
190 ASN O	-6.57	28.50	70.46	15.00
191 LYS N	-7.23	29.26	68.47	15.00
191 LYS CA	-7.79	30.49	68.99	15.00
191 LYS CB	-9.10	30.17	69.71	15.00
191 LYS CG	-10.01	29.28	68.87	15.00
191 LYS CD	-10.50	29.93	67.53	15.00
191 LYS CE	-9.41	30.19		15.00
191 LYS NZ	-8.63	28.95	66.03	15.00
191 LYS C	-6.81	31.27	69.84	15.00
191 LYS 0	-7.13	31.73	70.96	15.00
192 GLY N	-5.60	31.43	69.30	15.00
192 GLY CA	-4.58	32.18	70.01	15.00
192 GLY C	-3.63	31.36	70.87	
192 GLY 0	-2.54	31.86	71.19	
193 TYR N	-4.03	30.16	71.27	15.00
193 TYR CA	-3.20	29.32	72.12	15.00
193 TYR CB	-4.03	28.72	73.28	15.00
193 TYR CG	-4.53	29.79	74.21	15.00
193 TYR CD1	-5.69	30.50	73.91	15.00
193 TYR CE1	-6.13	31.53	74.73	15.00
193 TYR CD2	-3.81	30.13	75.36	15.00
193 TYR CE2	-4.22	31.14	76.20	15.00
193 TYR CZ	-5.37	31.84	75.89	15.00
193 TYR OH	-5.74	32.89	76.71	15.00
193 TYR C	-2.54	28.21	71.35	15.00
193 TYR O	-2.85	27.94	70.17	15.00
194 ILE N	-1.68		72.06	15.00
194 ILE CA	-0.99	26.35	71.50	15.00
194 ILE CB	0.26	26.72	70.64	15.00
194 ILE CG2	1.18	27.65	71.44	15.00
194 ILE CG1	1.03	25.47		15.00
194 ILE CD1 194 ILE C	2.16	25.76	69.14	15.00
	-0.54	25.54	72.70	15.00
194 ILE 0	-0.28	26.08	73.75	15.00
195 LEU N	-0.56	24.23	72.57	15.00
195 LEU CA	-0.12	23.30	73.59	15.00
195 LEU CB	-1.11	22.15	73.67	15.00
195 LEU CG	-2.34	22.29	74.58	15.00
195 LEU CD1	-2.88	23.70	74.67	15.00
195 LEU CD2	-3.41	21.38	74.05	15.00
195 LEU C	1.21	22.91	73.09	15.00
195 LEU 0	1.31	22.26	71.99	15.00

196 MET N	2.25	23.13	73.83	15.00
196 MET CA	3.58	22.69		
196 MET CB	4.57	23.83		
196 MET CG	4.29	24.99		
196 MET SD	5.56	26.26		
196 MET CE	7.08	25.68		
196 MET C	3.97	21.45		
196 MET 0	3.31	21.15		
197 ALA N	4.97	20.70		15.00
197 ALA CA	5.38	19.48		15.00
197 ALA CB	6.32	18.63	73.67	15.00
197 ALA C	6.01	19.71	75.90	15.00
197 ALA O	6.93	20.50		
198 ARG N	5.56	19.01		
198 ARG CA	6.13	19.21		
198 ARG CB	5.05	19.61	79.26	15.00
198 ARG CG	5.46	19.52	80.76	
198 ARG CD	4.45	20.21	81.65	
198 ARG NE	3.20	19.48	81.76	15.00
198 ARG CZ	3.00	18.51	82.64	15.00
198 ARG NH1	3.99	18.18	83.47	15.00
198 ARG NH2	1.80	17.95	82.78	15.00
198 ARG C	6.82	17.93	78.72	15.00
198 ARG O	6.19	16.88	78.82	15.00
199 ASN N	8.13		78.90	15.00
199 ASN CA	8.99		79.34	15.00
199 ASN CB	8.28	15.93	80.33	15.00
199 ASN CG	8.26	16.47	81.77	15.00
199 ASN OD1	7.22	16.83	82.31	15.00
199 ASN ND2	9.45	-	82.38	15.00
199 ASN C	9.79	16.12	78.28	15.00
199 ASN O	10.49	15.17	78.61	15.00
200 LYS N	9.77	16.62	77.05	15.00
200 LYS CA	10.56	16.06	75.94	15.00
200 LYS CB	9.83	16.23	74.61	15.00
200 LYS CG	8.82	15.14	74.28	15.00
200 LYS CD	7.48	15.29	74.99	15.00
200 LYS CE	6.27	15.00	74.07	15.00
200 LYS NZ	6.29	13.60	73.43	15.00
200 LYS C	11.87	16.86	75.92	15.00
200 LYS O	12.15	17.61	74.99	15.00
201. ASN N	12.64	16.70	76.99	15.00
201 ASN CA	13.91	17.41	77.22	15.00

201 ASN CB	15.14	16.71	76.60	15.00
201 ASN CG	15.48	15.43	77.29	15.00
201 ASN OD1	15.33	14.36	76.70	15.00
201 ASN ND2	15.91	15.52	78.56	
201 ASN C	13.91	18.86	76.81	
201 ASN 0	14.74	19.28	76.01	15.00
202 ASN N	13.02	19.62	77.41	15.00
202 ASN CA	12.87	21.05		
202 ASN CB	13.98	21.82		
202 ASN CG	13.66	23.27	78.07	15.00
202 ASN OD1	12.53	23.63	78.38	15.00
202 ASN ND2	14.63	24.12	77.83	15.00
202 ASN C	12.73	21.47	75.69	
202 ASN O	13.48	22.28		
203 ALA N	11.66	21.00		15.00
203 ALA H	11.06	20.52	75.65	15.00
203 ALA CA	11.34	21.22	73.64	15.00
203 ALA CB	10.03	20.53	73.27	
203 ALA C	11.16	22.72	73.38	
203 ALA O	10.25	23.38	73.78	15.00
204 CYS N	11.98	23.21	72.42	15.00
204 CYS CA	12.00	24.60	72.00	15.00
204 CYS C	12.56	25.58	73.01	15.00
204 CYS O	12.50	26.79	72.81	15.00
204 CYS CB	10.64		71.56	15.00
204 CYS SG	10.05	24.34	69.96	15.00
205 GLY N	13.03	25.04	74.13	15.00
205 GLY CA	13.63	25.86	75.14	15.00
205 GLY C	12.69	26.61	76.01	15.00
205 GLY O	13.03		76.63	15.00
206 ILE N	<b>1</b> 1.50	26.08	76.12	15.00
206 ILE CA	10.47	26.69	76.92	15.00
206 ILE CB	9.30	25.69	77.08	15.00
206 ILE CG2	9.78	24.44	77.78	15.00
206 ILE CG1	8.15	26.33	77.87	15.00
206 ILE CD1	7.20	27.12	77.00	15.00
206 ILE C	10.95	27.11	78.33	15.00
206 ILE O	10.50	28.12	78.86	15.00
207 ALA N	11.85	26.34	78.95	15.00
207 ALA CA	12.32	26.67	80.30	15.00
207 ALA CB	12.30	25.45	81.18	15.00
207 ALA C	13.68	27.35	80.43	15.00
207 ALA O	14.20	27.46	81.56	15.00

208 ASN N	14.21	27.84	79.31	15.00
208 ASN CA	15.51	28.52	79.26	
208 ASN CB	16.13		77.88	15.00
208 ASN CG	16.75		77.59	15.00
208 ASN OD1	16.26	25.96	78.00	15.00
208 ASN ND2	17.86	27.03	76.88	15.00
208 ASN C	15.37	30.02	79.55	
208 ASN O	16.37	30.73	79.73	
209 LEU N	14.16	30.55		
209 LEU CA	13.97	31.98	79.73	15.00
209 LEU CB	14.05	32.79	78.43	15.00
209 LEU CG	14.45	34.26	78.52	15.00
209 LEU CD1	15.95	34.30	78.78	
209 LEU CD2	14.12	35.06	77.23	15.00
209 LEU C	12.71	32.35	80.51	15.00
209 LEU O	12.13	33.44	80.31	15.00
210 ALA N	12.37	31.55	81.52	15.00
210 ALA H	12.79	30.67	81.51	15.00
210 ALA CA	11.14	31.81	82.27	15.00
210 ALA CB	10.61	30.53	82.92	15.00
210 ALA C	11.43	32.81	83.41	15.00
210 ALA O	12.41	32.78	84.11	15.00
211 SER N	10.44	33.74	83.58	15.00
211 SER CA	10.44	34.68	-	15.00
211 SER CB	11.25	35.95	84.38	15.00
211 SER OG	10.66	36.74	83.37	15.00
211 SER C	8.97	35.02	85.03	15.00
211 SER 0	8.07	34.77	84.22	15.00
212 PHE N	8.74	35.50	86.25	15.00
212 PHE CA	7.44	35.96	86.75	15.00
212 PHE CB	6.68	34.87	87.53	15.00
212 PHE CG	7.43	34.30	88.72	15.00
212 PHE CD1	8.34	33.31	88.54	15.00
212 PHE CD2	7.26	34.79	90.01	15.00
212 PHE CE1	9.08	32.83	89.64	15.00
212 PHE CE2	8.00	34.30	91.08	15.00
212 PHE CZ	8.90	33.33	90.90	15.00
212 PHE C	7.65	37.23	87.63	15.00
212 PHE O	8.64	37.35	88.34	15.00
213 PRO N	6.79	38.23	87.46	15.00
213 PRO CD	5.64	38.27	86.55	15.00
213 PRO CA	6.88	39.49	88.23	15.00
213 PRO CB	6.03	40.48	87.41	15.00

213	PRO	CG	4.96	39.59	87.00	15.00
213	PRO	C	6.30	39.32	89.66	15.00
213	PRO	0	5.25	38.68	89.88	15.00
214	LYS	N	6.99	39.82	90.67	15.00
214	LYS	CA	6.40	39.71	92.01	15.00
214	LYS	CB	7.46	39.54	93.11	15.00
214	LYS	CG	8.31	38.28	92.90	15.00
214	LYS	CD	8.81	37.72	94.19	15.00
214	LYS	CE	9.63	38.73	94.95	15.00
214	LYS	NZ	10.60	39.41	94.05	15.00
214	LYS	С	5.57	40.96	92.27	15.00
214			5.99	42.07	91.94	15.00
	MET		4.33	40.78	92.72	15.00
	MET		3.47	41.93	92.99	15.00
	MET	CB	2.11	41.74		15.00
	MET		1.57	43.08	91.85	15.00
	MET		0.31	43.10	90.56	15.00
	MET		-1.09	43.67	91.53	15.00
	MET		3.31	42.16	94.50	15.00
	MET		3.48	41.21		15.00
215		OT2	3.17	43.33	94.89	15.00
216		OH2	8.87	46.84	97.48	15.00
217		OH2	-2.18	37.97	73.56	15.00
218		OH2	1.71	36.04	75.21	15.00
219		OH2	9.44	52.65	61.91	15.00
220		OH2	0.80	56.90	67.17	15.00
221		OH2	-2.51	36.41	82.35	15.00
222	НОН		17.40	43.23	83.47	15.00
223	HOH		-1.57	52.44	64.46	15.00
22 <b>4</b> 225	HOH		12.41	35.91	80.62	15.00
225	НОН		11.65 11.38	62.93	58.36	15.00
227			5.00	48.93 12.95	74.41 78.69	15.00 15.00
	нон		4.86	15.66	86.17	15.00
	нон		-9.01	32.96	72.96	15.00
	нон		14.02	19.79	82.02	15.00
231	нон		18.09	36.59	88.86	15.00
	нон		0.22	37.62	76.69	15.00
	нон		3.45	36.52	73.19	15.00
	нон		13.53	38.17	80.00	15.00
		OH2		48.59	69.63	15.00
	нон		-5.38	44.85	97.00	15.00
	нон		-7.89	45.15	89.13	15.00
					~~. <b>~~</b>	

		H OH2	2.43	19.39	65.70	15.00
239		H OH2	7.43	21.65	71.07	15.00
240		н он2	2.41	16.41	85.78	15.00
241	l HOI	H OH2	-0.33	36.99	59.82	15.00
242	OH S	OH2	-7.54	26.54	72.89	15.00
243	HOP	OH2	-3.03	44.85	65.86	15.00
244	HOH	H OH2	0.80	35.69	86.28	15.00
245	HOH	OH2	-9.57	36.85	95.54	15.00
246	HOH	OH2	-1.06	62.37	83.93	15.00
247	HOH	OH2	7.72	62.09	69.81	15.00
248	HOH	OH2	4.95	60.85	80.90	15.00
249	HOH	OH2	14.51	30.92	83.13	15.00
250	HOH	OH2	-1.50	28.47	63.31	15.00
251	нон	OH2	15.32	22.32	71.31	15.00
252	HOH	OH2	-1.00	14.71	55.75	15.00
253	HOH	OH2	6.77	18.87	84.05	15.00
254			-9.65	32.88	79.27	15.00
255	HOH	OH2	-2.32	34.26	69.09	15.00
256	•	OH2	-11.12	32.20	64.94	15.00
257		OH2	-3.80	45.19	72.07	15.00
258		OH2	-7.43	38.35	65.10	15.00
259			1.41	46.77	63.08	15.00
260	HOH	OH2	-3.18	37.41	80.05	15.00
261	HOH	OH2	7.12	59.13	81.53	15.00
262	HOH	OH2	9.18	59.65	79.58	15.00
263	HOH	OH2	8.43	57.49	83.56	15.00
264	HOH	OH2	22.06	33.25	80.24	15.00
265	HOH	OH2	20.66	27.84	95.17	15.00
266	HOH	OH2	17.09	49.08	84.72	15.00
267	HOH	OH2	12.06	54.25	84.82	15.00
268		OH2	9.93	50.78	92.92	15.00
269		OH2	13.59	41.50	91.19	15.00
270		OH2	11.18	49.64	64.47	15.00
	нон		12.14	55.71	75.81	15.00
	HOH		9.07	26.37	66.15	15.00
273			24.27	24.31	64.11	15.00
274	НОН		18.35	21.16	79.19	15.00
	нон		20.62	28.49	61.87	15.00
	нон		13.58	15.19	72.83	15.00
277	HOH	OH2	9.33	19.74	77.14	15.00

#### TABLE XI

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
2420H2	25C1	180D1	69.66	2420H2	25C1	18ND2	48.33
2420H2	25C1	1840	82.25	2420H2	25C1	18CG	62.66
2420H2	25C1	184C	92.20	180D1	25C1	184CA	78.01
180D1	25C1	18ND2	33.84	180D1	25C1	1840	82.68
180D1	25C1	184CD1	97.11	180D1	25C1	18CG	16.91
180D1	25C1	184C	73.04	180D1	25C1	200	60.56
184CB	25C1	184CA	23.31	184CB	25C1	18ND2	93.14
184CB	25C1	1840	42.27	184CB	25C1	184CG	22.32
184CB	25C1	184CD1	38.40	184CB	25C1	18CG	92.58
184CB	25C1	184C	36.61	184CB	25C1	184CD2	32.79
184CB	25C1	184NE1	49.18	184CA	25C1	18ND2	72.65
184CA	25C1	1840	35.91	184CA	25C1	184CG	38.24
184CA	25C1	184CD1	44.79	184CA	25C1	18CG	69.61
184CA	25C1	184C	21.48	184CA	25C1	184CD2	52.43
184CA	25C1	184NE1	58.86	18ND2	25C1	1840	58.28
18ND2	25C1	18CG	18.49	18ND2	25C1	184C	56.82
18ND2	25C1	200	92.88	1840	25C1	184CG	64.55
1840	25C1	184CD1	78.02	1840	25C1	18CG	67.10
1840	25C1	184C	17.17	1840	25C1	184CD2	73.76
1840	25C1	184NE1	90.68	18 <b>4CG</b>	25C1	184CD1	19.59
184CG	25C1	184C	56.91	184CG	25C1	184CD2	15.07
184CG	25C1	184NE1	27.32	184CD1	25C1	18CG	99.42
184CD1	25C1	184C	66.13	184CD1	25C1	184CD2	27.32
184CD1	25C1	184NE1	14.36	184CD1	25C1	200	86.22
18CG	25C1	184C	59.85	18CG	25C1	200	77.46
184C	25C1	184CD2	69.30	184C	25C1	184NE1	80.01
184CD2	25C1	184NE1	26.78	184NE1	25C1	200	80.98
180D1	25C2	184CA	93.13	1.80 <b>D1</b>	25C2	18CG	18.16
180D1	25C2	20N	42.62	180D1	25C2	18ND2	35.44
180D1	25C2	2420H2	68.83	180D1	25C2	200	79.33
180D1	25C2	1830	73.73	180 <b>D</b> 1.	25C2	184C	80.63
180D1	25C2	19CG	85.64	180D1	25C2	20CA	45.72
180D1	25C2	19N	47.90	180 <b>D1</b>	25C2	1840	85.52
180D1	25C2	20C	65.28	18001	25C2	184N	90.71
180D1	25C2	19C	49.49	180D1	25C2	18CB	22.83

			•	TABLE XI			
180D1		· -	82.72		. 25C2	18CA	34.96
180D1			34.23	180D1	25C2		54.26
184CD1			49.90	184CD1	25C2		40.27
184CD1	25C2		20.32	184CD1	25C2		55.01
184CD1	25C2	184C	70.22	184CD1			61.53
184CD1	25C2	19N	85.18	184CD1		1840	78.11
184CD1	25C2	184NE1	15.62	184CD1		184N	44.48
184CD1	25C2	183C	47.20	184CD1	25C2	18CA	94.36
184CD1	25C2	18C	95.90	184CD1	25C2	184CD2	23.77
184CD1	25C2	19CA	87.00	184CA	25C2	18CG	78.59
184CA	25C2	18ND2	76.16	184CA	25C2	184CB	23.25
184CA	25C2	184CG	39.72	184CA	25C2	1830	40.92
184CA	25C2	184C	20.77	184CA	25C2	19CG	86.38
184CA	25C2	19N	76.16	184CA	25C2	1840	33.25
184CA	25C2	184NE1	65.31	184CA	25C2	184N	13.46
184CA	25C2	18CB	70.34	184CA	25C2	183C	28.93
184CA	25C2	18CA	61.99	184CA	25C2	18C	75.30
184CA	25C2	184CD2	51.97	184CA	25C2	19CA	89.58
18CG	25C2	20N	59.81	18CG	25C2	18ND2	20.07
18CG	25C2	184CB	99.39	18CG	25C2	2420H2	62.04
18CG	25C2	200	97.48	18CG	25C2	1830	69.32
18CG	25C2	184C	63.66	18CG	25C2	19CG	95.51
18CG	25C2	20CA	63.78	18CG	25C2	19N	54.69
18CG	25C2	1840	67.41	18CG	25C2	20C	83.28
18CG	25C2	184N	79.10	18CG	25C2	19C	65.07
18CG	25C2	18CB	12.54	18CG	25C2	183C	75.08
18CG	25C2	18CA	30.58	18CG	25C2	18C	39.48
18CG	25C2	19CA	65.54	20N	25C2	18ND2	78.04
20N	25C2	200	42.02	20N	25C2	1830	74.98
20N	25C2	1.9CG	52.80	20N	25C2	20CA	19.88
20N	25C2	19N	37.71	20N	25C2	20C	32.71
20N	25C2	19C	11.44	2 <b>0N</b>	25C2	18CB	58.04
20N	25C2	183C	87.83	20 <b>N</b>	25C2	18CA	55.70
20N	25C2	18C	39.13	2 <b>0N</b>	25C2	19CA	28.06
	25C2	184CB	92.26	18ND2	25C2	2420H2	45.09
	25C2	1830	90.83	18ND2	25C2	184C	57.03
	25C2	20CA	77.75	18ND2	25C2	19N	73.94
	25C2	1840	55.05	18ND2	25C2	20C	96.02
		134N	81.33	18ND2	25C2	19C	84.52
	25C2	18CB	29.48	1 & NID2	25C2	183C	82.74
	25C2	18CA	46.57	18ND2	25C2	18C	58.80
	25C2	19CA	85.55	184CB	25C2	184CG	22.40
		1830		184CB	25C2	184C	36.03
184CB	25C2	19CG	94.64	184CB	25C2	19 <b>N</b>	96.61

			т	ABLE XI			
184CB	25C2	1840	38.77	184CB	25C2	184NE1	52.96
184CB	25C2	184N	31.46	184CB	25C2	18CB	92.63
184CB	25C2	183C	46.49	184CB	25C2	18CA	85.24
184CB	25C2	18C	97.95	184CB	25C2	184CD2	31.97
2420H2	25C2	184C	81.46	2420H2	25C2	20CA	87.88
2420H2	25C2	1840	68.50	2420H2	25C2	20C	95.83
2420H2	25C2	18CB	73.69	2420H2	25 <b>C</b> 2	18CA	91.46
184CG	25C2	1830	61.89	184CG	25C2	184C	57.17
184CG	25C2	19CG	81.07	184CG	25C2	19N	97.59
184CG	25C2	1840	61.14	184CG	25C2	184NE1	30.80
184CG	25C2	184N	40.66	184CG	25C2	183C	50.54
184CG	25C2	18CA	97.12	184CG	25C2	184CD2	12.85
200	25C2	19CG	59.65	200	25C2	20CA	33.80
200	25C2	19N	73.93	200	25C2	184NE1	98.68
200	25C2	20C	15.00	200	25C2	19C	44.20
200	25C2	18CB	98.94	200	25C2	18CA	97.51
200	25C2	18C	80.19	200	25C2	19CA	58.60
1830	25C2	184C	52.11	1830	25C2	19CG	50.15
1830	25C2	20CA	94.61	1830	25C2	19N	37.44
1830	25C2	1840	68.46	1830	25C2	184NE1	67.20
1830	25C2	184N	28.51	1830	25C2	19C	66.39
1830	25C2	18CB	56.86	1830	25C2	183C	12.86
1830	25C2	18CA	39.61	1830	25C2	18C	42.74
1830	25C2	184CD2	73.55	1830	25C2	19CA	48.97
184C	25C2	19N	79.88	184C	25C2	1840	16.37
184C	25C2	184NE1	85.33	184C	25C2	184N	31.43
184C	25C2	18CB	58.56	184C	25C2	183C	43.26
184C	25C2	18CA	56.66	184C	25C2	18C	73.56
184C	25C2	184CD2	67.95	184C	25C2	19CA	95.56
19CG	25C2	20CA	68.54	19CG	25C2	19N	41.00
19CG	25C2	184NE1	59.71	19CG	25C2	20C	65.93
19CG	25C2	184N	72.99	19CG	25C2	19C	41.61
19CG	25C2	18CB	85.55	19CG	25C2	183C	58.81
19CG	25C2	18CA	69.73	19CG	25 <b>C2</b>	18C	56.16
19CG	25C2	184CD2	84.49	19CG	25C2	19CA	31.40
20CA	25C2	19N	57.18	20CA	25C2	20C	19.60
20CA	25C2	19C	30.27	20CA	25C2	18CB	66.65
20CA	25C2	1.8CA	70.02	20CA	25C2	18C	55.87
20CA	25C2	19CA	47.75	19N	25C2	1840	94.75
19N	25C2	184NE1	91.69	19N	25C2	20C	69.08
19N	25C2	184N	65.26	19N	25C2	19C	30.47
19N	25C2	1.8CB	44.60	19N	25C2	183C	50.23
19N	25C2	18CA	30.19	19N	25C2	18C	15.21
19N	25C2	19CA	16.11	1840	25C2	184NE1	91.68

			T	ABLE XI			
1840	25C2	184N	45.89	1840	25C2	18CB	65.91
1840	25C2	183C	59.15	1840	25C2	18CA	68.65
1840	25C2	18C	86.43	1840	25C2	184CD2	69.23
184NE1	25C2	184N	60.01	184NE1	25C2	183C	61.27
184NE1	25C2	184CD2	26.96	184NE1	25C2	19CA	89.32
20C	25C2	19C	38.71	20C	25C2	18CB	86.12
20C	25C2	18CA	87.75	20C	25C2	18C	71.80
20C	25C2	19CA	55.73	184N	25C2	19C	94.86
184N	25C2	18CB	68.81	184N	25C2	183C	15.92
184N	25C2	18CA	56.71	184N	25C2	18C	67.03
184N	25C2	184CD2	53.50	184N	25C2	19CA	77.48
19C	25C2	18CB	60.71	19C	25C2	183C	79.19
19C	25C2	18CA	54.22	19C	25C2	18C	36.38
19C	25C2	19CA	17.76	18CB	25C2	183C	63.21
18CB	25C2	18CA	18.13	18CB	25C2	18C	29.64
18CB	25C2	19CA	57.42	183C	25C2	18CA	47.76
183C	25C2	18C	54.26	183C	25C2	184CD2	62.80
183C	25C2	19 <b>CA</b>	61.67	18CA	25C2	18C	17.97
18CA	25C2	19CA	45.40	18C	25C2	19CA	28.26
200	25C3	19CG	77.70	200	25C3	20N	48.86
200	25C3	180D1	82.41	200	25C3	20C	15.02
200	25C3	20CA	36.52	200	25C3	19CD	82.71
200	25C3	19C	49.69	200	25C3	19N	81.60
200	25C3	19CB	69.52	200	25C3	18CG	93.56
200	25C3	190E1	97.10	200	25C3	19CA	66.55
184CD1	25C3	19CG	70.95	184CD1	25C3	184NE1	20.32
184CD1	25C3	184CG	17.18	184CD1	25C3	19CD	63.15
184CD1	25C3	19N	84.37	184CD1	25C3	184CA	41.74
184CD1	25C3	1830	51.08	184CD1	25C3	19CB	82.21
184CD1	25C3	184CB	33.05	184CD1	25C3	18CG	99.94
184CD1	25C3	190E1	48.60	184CD1	25C3	19CA	92.55
184CD1	25C3	184CE2	24.46	19CG	25C3	20N	60.87
19CG	25C3	180D1	86.79	19CG	25C3	20C	80.83
19CG	25C3	184NE1	70.80	19CG	25C3	20CA	78.71
19CG	25C3	184CG	86.74	19CG	25C3	19CD	19.15
19CG	25C3	19C	46.56	19CG	25C3	19N	42.15
19CG	25C3	184CA	83.50	19CG	25C3	1830	50.42
19CG	25C3	19CB	14.34	19CG	25C3	184CB	93.55
19CG	25C3	18CG	90.03	19CG	25C3	190E1	29.89
19CG	25C3	19CA	32.74	19CG	25 <b>C3</b>	184CE2	83.95
20N	25C3	180D1	40.95	20N	25C3	20C	38.02
20N	25C3	20CA	21.15	20 <b>N</b>	25C3	19CD	78.19
2GN	25C3	19C	14.72	20N	25C3	19N	36.90
20N	25C3	184CA	98.63	20N	25C3	1830	71.83

			7	ABLE XI			
20N	25C3	19CB	46.57	20N	25C3	18CG	51.19
20N	25C3	190E1	90.75	20N	25C3	19CA	30.28
20N	25C3	2420H2	86.06	180D1	25C3	20C	67.75
180D1	25C3	20CA	45.89	180D1	25C3	19C	52.30
180D1	25C3	19N	45.18	180D1	25C3	184CA	70.04
180D1	25C3	1830	62.55	180D1	25C3	19CB	74.34
180D1	25C3	184CB	86.92	180D1	25C3	18CG	11.19
180D1	25C3	19CA	55.77	180D1	25C3	2420H2	51.35
20C	25C3	20CA	22.11	20C	25C3	19CD	90.19
20C	25C3	19C	43.18	20C	25C3	19N	73.89
20C	25C3	19CB	69.80	20C	25C3	18CG	78.83
20C	25C3	19CA	61.77	20C	25C3	2420H2	89.06
184NE1	25C3	184CG	32.04	184NE1	25C3	19CD	56.99
184NE1	25C3	19N	96.67	184NE1	25C3	18 <b>4CA</b>	61.92
184NE1	25C3	1830	67.54	184NE1	25C3	19CB	84.48
184NE1	25C3	184CB	50.65	184NE1	25 <b>C</b> 3	190E1	42.30
184NE1	25C3	19CA	99.57	184NE1	25C3	184CE2	13.16
20CA	25C3	19CD	93.68	20CA	25C3	19C	32.85
20CA	25C3	19N	57.68	20CA	25C3	1830	91.89
20CA	25C3	19CB	64.99	20CA	25C3	18CG	57.06
20CA	25C3	19CA	50.80	20CA	25C3	2420H2	76.89
184CG	25C3	19CD	80.27	184CG	25C3	19N	91.72
184CG	25C3	184CA	34.06	184CG	25C3	1830	56.45
184CG	25C3	19CB	96.63	184CG	25C3	184CB	18.84
184CG	25C3	18CG	93.58	184CG	25C3	190E1	65.78
184CG	25C3	184CE2	28.26	184CG	25C3	2420H2	94.95
19CD	25C3	19C	63.50	19CD	25C3	19N	60.85
19CD	25C3	184CA	88.20	19CD	25C3	1830	60.91
19CD	25C3	19CB	32.82	19CD	25C3	184CB	92.35
19CD	25C3	190E1	14.96	19CD	25C3	19CA	51.66
19CD	25C3	184CE2	69.81	19C	25C3	19N	31.91
19C	25C3	184CA	98.36	19C	25C3	1830	66.62
19C	25C3	19CB	32.43	19C	25C3	18CG	61.26
19C	25C3	190E1	76.35	1.9C	25C3	19CA	18.84
19N	25C3	184CA	66.70	19N	25C3	1830	35.29
19N	25C3	19CB	31.75	19N	25C3	184CB	85.03
19N	25C3	18CG	47.94	19N	25C3	190E1	67.18
19N	25C3	19CA	17.97	19N	25C3	2420H2	95.18
184CA	25C3	1830	35.33	184CA	25 <b>C</b> 3	19CB	86.05
184CA	25C3	184CB	19.26	184CA	25C3	18CG	59.77
184CA	25C3	190E1	78.16	184CA	25C3	19CA	83.25
184CA	25C3	184CE2	61.84	184CA	25C3	2420H2	75.15
1830	25C3	19CB	50.84	183C	25C3	184CB	51.32
1830	25C3	18CG	57.48	1830	25C3	190E1	57.08

			7	ABLE XI			
1830	25C3	19CA	49.31	1830	25C3	184CE2	75.42
1830	25C3	2420H2	96.01	19CB	25C3	184CB	99.69
19CB	25C3	18CG	79.10	19CB	25C3	190E1	44.22
19CB	25C3	19CA	18.92	19CB	25C3	184CE2	97.58
184CB	25C3	18CG	76.03	184CB	25C3	190E1	79.22
184CB	25C3	184CE2	46.73	184CB	25C3	2420H2	78.05
18CG	25C3	19CA	61.53	18CG	25C3	2420H2	47.27
190 <b>E</b> 1	25C3	19CA	62.05	190E1	25C3	184CE2	55.29
200	25C4	20C	10.43	200	25C4	19CG	62.30
200	25C4	20N	35.25	200	25C4	20CA	26.32
200	25C4	19CD	71.57	200	25C4	21NE2	50.82
200	25C4	180D1	61.89	184CD1	25C4	184NE1	20.27
184CD1	25C4	19CG	57.56	184CD1	25C4	184CG	16.41
184CD1	25C4	184CE2	28.68	184CD1	25C4	20N	90.35
184CD1	25C4	19CD	54.84	184CD1	25C4	180D1	81.65
184CD1	25C4	184CD2	25.56	184NE1	25C4	19CG	61.65
184NE1	25C4	184CG	29.79	184NE1	25C4	184CE2	15.87
184NE1	25C4	19CD	51.78	184NE1	25C4	184CD2	25.93
20C	25C4	19CG	66.50	20C	25C4	20N	31.03
20C	25 <b>C4</b>	20CA	17.90	20C	25C4	19CD	78.21
20C	25C4	21NE2	44.60	20C	25C4	180D1	53.90
19CG	25C4	184CG	72.66	19CG	25C4	184CE2	77.52
19CG	25C4	20N	46.19	19CG	25C4	20CA	61.94
19CG	25C4	19CD	18.17	19CG	25C4	180D1	63.67
19CG	25C4	184CD2	82.38	184CG	25C4	184CE2	28.98
184CG	25C4	20N	98.65	184CG	25C4	19CD	71.23
184CG	25C4	180D1	82.35	184CG	25C4	184CD2	16.48
184CE2	25C4	19CD	67.12	184CE2	25C4	184CD2	16.42
20N	25C4	20CA	17.47	2 <b>0N</b>	25C4	19CD	63.14
20N	25C4	21NE2	68.18	2 <b>0N</b>	25C4	180D1	30.69
20CA	25C4	19CD	77.50	20CA	25C4	21NE2	50.79
20CA	25C4	180D1	36.03	19CD	25 <b>C4</b>	180D1	81.70
19CD	25C4	184CD2	76.27	21 <b>NE2</b>	25C4	180D1	71.09
180D1	25C4	184CD2	98.63	18 <b>4CD1</b>	25C5	200	89.99
184CD1	25C5	184NE1	17.97	184CD1	25C5	184CG	17.29
184CD1	25C5	184CE2	27.88	184CD1	25C5	2420H2	99.26
1.84CD1	25C5	184CD2	27.37	200	25C5	184NE1	92.08
200	25C5	2420H2	86.50	200	25C5	21NE2	47.88
184NE1	25C5	184CG	28.79	184NE1	25C5	184CE2	16.62
184NE1	25C5	184CD2	27.40	184CG	25C5	184CE2	28.30
184CG	25C5	2420H2	92.08	184CG	25C5	184CD2	17.13
184CE2	25C5	184CD2	16.84	2420H2	25C5	21NE2	65.12
2420H2	25C6	184CB	96.30	2420H2	25C6	184CA	83.88
2420H2	25C6	180D1	51.77	2420H2	25C6	1840	63.15

			•	TABLE XI			
184CG	25C6		17.92		25C6	184CB	19.89
184CG	25C6		17.11	184CG	25C6	184NE1	27.45
184CG	25C6	184CA	31.62	184CG	25C6		84.34
184CG	25C6		53.51	184CD1	. 25C6		33.90
184CD1		184CD2	27.70	184CD1			16.02
184CD1	25C6	184CA	37.26				76.75
184CD1	25C6	1840	64.28	184CB	25C6		32.92
184CB	25C6	184NE1	46.79	184CB	25C6		18.16
184CB	25C6	180D1	76.20	184CB	25C6		33.78
184CD2	25C6	184NE1	26.79	184CD2		_ <del>_</del>	47.98
184CD2	25C6	1840	65.87	184NE1	25C6		53.03
184NE1	25C6	180D1	88.79	184NE1	25C6	1840	79.21
184CA	25C6	180D1	58.04	184CA	25C6	1840	28.15
180D1	25C6	1840	61.50	200	25C7	20C	4.43
200	25C7	19CG	62.28	200	25C7	19CD	77.38
200	25C7	19NE2	74.82	200	25C7	190E1	90.67
184NE1	25C7	19CG	59.54	184NE1	25C7	19CD	53.46
184NE1	25C7	184CD1	17.49	184NE1	25C7	19NE2	66.80
184NE1	25C7	190E1	40.08	184NE1	25C7	184CE2	14.60
20C	25C7	19CG	65.46	20C	25C7	19CD	81.18
20C	25C7	19NE2	79.11	20C	25C7	190E1	94.22
19CG	25C7	19CD	19.72	19CG	25C7	184CD1	52.93
19CG	25C7	19NE2	30.67	19CG	25C7	190E1	29.47
19CG	25C7	184CE2	74.12	19CD	25 <b>C</b> 7	184CD1	53.97
19CD	25C7	19NE2	16.35	19CD	25C7	190E1	14.31
19CD	25C7	184CE2	67.37	184CD1	25C7	19NE2	69.74
184CD1	25C7	190E1	43.73	184CD1	25 <b>C</b> 7	184CE2	26.87
19NE2	25C7	190E1	26.90	19NE2	25C7	184CE2	79.58
190E1	25C7	184CE2	53.50	184NE1	2508	19CD	69.28
184NE1	2508	19NE2	88.26	184NE1	2508	190E1	52.70
184NE1	2508	19CG	72.15	134NE1	2508	184CD1	18.62
184NE1	2508	184CE2	15.78	184NE1	2508	184CZ2	30.82
19CD	2508	19NE2	21.45	19CD	2508	190E1	19.40
19CD	2508	19CG	23.73	19CD	2508	200	83.60
19CD	2508	184CD1	65.36	19CD	2508	184CE2	84.12
19CD	2508	220	53.07	19CD	2508	184CZ2	93.15
19NE2	2508	190E1	35.59	19NE2	2508	19CG	38.13
19NE2	2508	200	82.17	19NE2	2508	184CD1	86.56
19NE2	2508	220	36.04	190E1	2508	19CG	37.46
190E1	2508	184CD1	53.61	1.90E1	2508	184CE2	66.40
190E1	2508	220	70.80	1.90E1	2508	184CZ2	73.99
19CG	2508	200	63.71	19CG	2508	184CD1	60.70
19CG	2508	184CE2	87.85	1.9CG	2508	220	54.69
200	2508	220	57.31	194CD1	2508	184CE2	30.67

			T	ABLE XI			
184CD1	2508	184CZ2	47.47	184CE2	2508	184CZ2	16.93
19NE2	25C9	184NE1	78.75	19NE2	25C9	19CD	19.52
19NE2	25C9	190E1	33.07	19NE2	25C9	184CE2	94.10
19NE2	25 <b>C</b> 9	19CG	30.70	19NE2	25C9	184CD1	73.44
19NE2	25C9	220	33.46	184NE1	25C9	19CD	59.66
184NE1	25C9	190E1	47.23	184NE1	25C9	184CE2	16.38
184NE1	25C9	184CZ2	33.08	184NE1	25C9	19CG	58.24
184NE1	25C9	184CD1	12.56	19CD	25C9	190E1	17.75
19CD	25C9	184CE2	75.46	19CD	25C9	184CZ2	89.07
19CD	25C9	19CG	17.50	19CD	25C9	184CD1	53.92
19CD	25C9	220	48.71	190E1	25C9	184CE2	61.55
190E1	25C9	184CZ2	73.08	190E1	25C9	19CG	30.61
190E1	25C9	184CD1	45.28	190E1	25C9	220	65.42
184CE2	25C9	184CZ2	17.56	184CE2	25C9	19CG	74.59
184CE2	25C9	184CD1	26.94	184CZ2	25C9	19CG	90.93
184CZ2	25C9	184CD1	44.43	19CG	25C9	184CD1	48.87
19CG	25C9	220	48.21	184CD1	25 <b>C</b> 9	220	97.02
19NE2	25010	23CA	53.09	19NE2	25010	19CD	16.71
19NE2	25010	220	36.83	19NE2	25010	23N	57.66
19NE2	25010	190E1	28.25	19 <b>NE</b> 2	25010	22C	50.13
23CA	25010	19CD	69.80	23CA	25010	220	36.05
23CA	25010	23N	17.68	23CA	25010	190E1	79.02
23CA	25010	22C	29.83	19CD	25010	220	49.55
19CD	25010	23N	73.42	19CD	25010	190E1	15.11
19CD	25010	22C	63.87	220	25010	23N	27.96
220	25010	190E1	63.96	220	25010	22C	14.56
23N	25010	190E1	85.79	23 <b>N</b>	25010	22C	15.87
190E1	25010	22C	77.94	162ND1	25C11	184CZ2	63.39
162ND1	25C11	162CE1	16.78	162ND1	25C11	184NE1	61.01
162ND1	25C11	184CE2	62.72	162ND1	25C11	162CG	15.67
162ND1	25C11	184CH2	68.93	162ND1	25C11	162CB	30.11
184CZ2	25C11	162CE1	53.21	184CZ2	25C11	184NE1	33.69
184CZ2	· 25C11	184CE2	16.65	184CZ2	25C11	162CG	59.25
184CZ2		184CH2	12.22	184CZ2		162CB	70.77
162CE1		184NE1	44.59	162CE1		184CE2	48.65
162CE1		162CG	27.93	162CE1		184CH2	61.58
162CE1		162CB	44.93	184NE1	25C11	184CE2	17.24
184NE1		162CG	67.01	184NE1		184CH2	45.73
184NE1		162CB	83.92	184CE2	25C11	162CG	63.63
184CE2		184CH2	28.52	184CE2	25C11	162CB	78.45
162CG		184CH2	61.52	162CG		162CB	17.47
184CH2		162CB	69.67	1380G		138CB	12.25
1380G		138CA	28.90	1380G		1610D1	38.49
138CB	25C15	138CA	18.42	138CB	25C15	1610D1	45.39

			T.	ABLE XI			
138CA	25C15	1610D1	48.24	162ND1	25C16	1610	83.50
162ND1	25C16	162CG	18.14	162ND1	25C16	162CE1	16.51
162ND1	25C16	162CB	37.71	162ND1	25C16	25 <b>S</b> G	53.12
162ND1	25C16	162CA	45.79	162ND1	25C16	161C	76.59
162ND1	25C16	184CZ2	58.80	162ND1	25 <b>C</b> 16	25CB	44.47
162ND1	25C16	162N	61.56	162ND1	25C16	190E1	54.44
1610	25C16	162CG	73.87	1610	25C16	162CE1	99.61
1610	25C16	162CB	55.47	1610	25C16	25SG	68.22
1610	25C16	162CA	38.54	1610	25C16	161C	12.03
1610	25C16	25CB	87.95	1610	25C16	162N	25.97
162CG	25C16	162CE1	32.06	162CG	25C16	162CB	21.17
162CG	25C16	25 <i>S</i> G	64.52	162CG	25C16	162CA	35.58
162CG	25C16	161C	64.63	162CG	25C16	184CZ2	58.11
162CG	25C16	25CB	60.75	162CG	25C16	162N	48.98
162CG	25C16	190E1	71.80	162CE1	25C16	162CB	53.00
162CE1	25C16	25 <i>S</i> G	57.54	162CE1	25C16	162CA	62.27
162CE1	25C16	161C	93.08	162CE1	25C16	184CZ2	50.19
162CE1	25C16	25CB	42.26	162CE1	25C16	162N	78.06
162CE1	25C16	190E1	40.00	162CB	25C16	25 <b>S</b> G	70.05
162CB	25C16	162CA	20.87	162CB	25C16	161C	44.96
162CB	25C16	184CZ2	71.73	162CB	25C16	25CB	73.86
162CB	25C16	162N	29.56	162CB	25C16	190E1	92.15
25SG	25C16	162CA	56.57	25SG	25C16	161C	72.16
25SG	25C16	25CB	21.35	25 <b>SG</b>	25C16	162N	67.66
25 <i>S</i> G	25C16	190E1	58.28	162CA	25C16	161C	30.81
162CA	25C16	184CZ2	91.78	162CA	25C16	25CB	67.26
162CA	25C16	162N	16.41	162CA	25C16	190E1	95.96
161C	25C16	25CB	89.77	161C	25C16	162N	15.65
184CZ2	25C16	25CB	89.70	184CZ2	25C16	190E1	64.53
25CB	25C16	162N	81.65	25CB	25C16	190E1	37.30
162ND1	25017	162CB	53.19	162ND1	25017	162CG	26.33
162ND1	25017	162CA	63.48	162ND1	25017	162N	85.18
162ND1	25017	162CE1	12.84	162ND1	25017	25 <b>S</b> G	56.41
162ND1	25017	162CD2	24.16	162ND1	25017	162C	54.50
162ND1	25017	162NE2	14.17	162ND1	25017	184CZ2	59.71
162ND1	25017	163N	47.86	162ND1	25017	25CB	41.51
162CB	25017	1610	76.56	162CB	25017	162CG	28.45
162CB	25017	162CA	28.85	162CB	25017	161C	60.68
162CB	25017	162N	40.36	162CB	25017	162CE1	64.12
162CB	25017	25 <i>S</i> G	83.81	162CB	25017	1610D1	58.91
162CB	25017	1.62CD2	36.23	162CB	25017	162C	31.62
162CB	25017	162NE2	<b>52.8</b> 3	162CB	25017	184CZ2	81.36
162CB	25017	163N	44.40	1.62CB	25017	25CB	83.22
152CB	25017	161CA	66.51	162CB	25017	161CB	74.63

			т	ABLE XI			
1610	25017	7 162CA	52.04	1610	25017	161C	18.10
1610	25017	7 162N	36.49	1610	25017	25SG	77.20
1610	25017	1610D1	60.19	1610	25017	162C	58.32
1610	25017	163N	63.78	1610	25017	25CB	97.03
1610	25017	161CA	21.34	1610	25017	161CB	35.39
162CG	25017	162CA	48.03	162CG	25017	161C	86.77
162CG	25017	162N	66.31	162CG	25017	162CE1	35.94
162CG	25017	25SG	73.55	162CG	25017	1610D1	84.63
162CG	25017	162CD2	9.57	162CG	25017		43.30
162CG	25017	162NE2	24.46	162CG	25017	184CZ2	61.90
162CG	25017	163N	46.12	162CG	25017	25CB	64.34
162CG	25017	161CA	94.08	162CA	25017	161C	40.53
162CA	25017	162N	22.22	162CA	25017	162CE1	76.31
162CA	25017		66.07	162CA	25017	1610D1	63.93
162CA	25017		57.57	162CA	25017	162C	11.05
162CA	25017		70.12	162CA	25017	163N	26.62
162CA	25017		74.60	162CA	25017	161CA	50.08
162CA	25017		64.72	161C	25017	162N	20.53
161C	25017		84.94	161C	25017	1610D1	45.99
161C	25017		95.86	161C	25017	162C	49.50
161C	25017		59.81	161C	25017		11.07
161C	25017	161CB	28.61	162N	25017	162CE1	97.92
162N	25017	25SG	81.29	162N	25017	1610D1	46.28
162N	25017	162CD2	75.34	162N	25017	162C	32.79
162N	25017	162NE2	90.12	162N	25017	163N	46.59
162N	25017	25CB	94.05	162N	25017	161CA	28.47
162N	25017	161CB	42.50	162CE1	25017	25SG	60.41
162CE1	25017	162CD2	30.63	162CE1	25017	162C	67.29
162CE1	25017	162NE2	14.05	162CE1	25017	184CZ2	50.87
162CE1	25017	163N	<b>59.7</b> 3	162CE1	25017	25CB	42.09
25SG	25017	162CD2	77.38	25SG	25017	162C	56.56
25SG 25SG	25017 25017	162NE2 25CB	70.26 20.52	25 <b>S</b> G	25017	163N	40.80
1610D1			88.85	25SG	25017	161CA	94.97
1610D1	25017	162CD2 163N	90.55	1610D1	25017		74.74
1610D1		161CB	28.90	1610D1		161CA	38.88
1610D1	25017		17.33	162CD2	25017		52.76
162CD2	25017	163N	54.44			184CZ2	52.52
162CD2	25017		62.89	162CD2	25017	25CB	65.18
162C	25017	1.62NE2 25CB	63.70	162C	25017 25017	163N	16.06
1.62C	25017	161CB	75.18	162C 162NE2			59.72
162NE2	25017	163N	59.43	162NE2		184CZ2 25CB	46.51
184CZ2	25017	25CB	88.90	163N	25017 25017		53.98
163N	25017	25CB 161CA	70.75	163N		25CB 161CB	48.07
~ ^ JM	んいひより	TOTCW	, , .	T0214	2 JUL /	TOTCD	87.59

		т	ABLE XI		
161CA	25017 161CB	17.64	25SG	25N18 162ND1	54.05
25SG	25N18 1610	72.12	25SG	25N18 25CB	22.85
25SG	25N18 19NE2	68.04	25SG	25N18 23CA	83.69
25SG	25N18 162CE1	55.83	25SG	25N18 162CA	54.53
25SG	25N18 19OE1	61.94	25SG	25N18 162CG	59.80
25 <i>S</i> G	25N18 162CB	64.56	162ND1	25N18 1610	73.75
162ND1	25N18 25CB	47.82	162ND1	25N18 19NE2	80.11
162ND1	25N18 162CE1	14.94	162ND1	25N18 162CA	39.62
162ND1	25N18 19OE1	53.76	162ND1	25N18 162CG	12.62
162ND1	25N18 162CB	29.54	1610	25N18 25CB	91.76
1610	25N18 162CE1	88.51	1610	25N18 162CA	34.27
1610	25N18 162CG	63.44	1610	25N18 162CB	47.06
25CB	25N18 19NE2	48.54	25CB	25N18 23CA	78.61
25CB	25N18 162CE1	42.68	25CB	25N18 162CA	66.30
25CB	25N18 19OE1	39.11	25CB	25N18 162CG	58.13
25CB	25N18 162CB	69.62	19NE2	25N18 23CA	48.32
19 <b>NE</b> 2	25N18 162CE1	66.29	19NE2	25N18 19OE1	27.62
19NE2	25N18 162CG	92.70	23CA	25N18 19OE1	75.78
162CE1	25N18 162CA	54.27	162CE1	25N18 19OE1	39.27
162CE1	25N18 162CG	26.90	162CE1	25N18 162CB	44.19
162CA	25N18 19OE1	90.56	162CA	25N18 162CG	30.46
162CA	25N18 162CB	17.73	190E1	25N18 162CG	66.12
190E1	25N18 162CB	83.30	162CG	25N18 162CB	17.35
25SG	25C19 1610	94.77	25 <b>S</b> G	25C19 162ND1	55.31
25SG	25C19 25CB	20.96	25 <b>S</b> G	25C19 162CA	63.69
25SG	25C19 161C	89.52	25 <b>S</b> G	25C19 23CA	94.30
25SG	25C19 23O	76.83	25 <b>S</b> G	25C19 23C	78.19
25SG	25C19 162N	77.13	25 <b>S</b> G	25C19 25N	39.34
25SG	25C19 19NE2	66.56	25 <b>S</b> G	25C19 163N	35.68
25 <i>S</i> G	25C19 162CE1	52.07	25SG	25C19 162CB	68.67
1610	25C19 162ND1	77.82	1610	25C19 162CA	38.27
1610	25C19 161C	6.45	1610	25C19 162N	21.62
1610	25C19 163N	59.92	1610	25C19 162CE1	90.20
				25C19 25CB	
	25C19 162CA	42.28		25C19 161C	71.68
	25C19 162N	57.38	162ND1		77.84
	25C19 19NE2		162ND1		45.25
162ND1	25C19 162CE1		162ND1		30.12
25CB	25C19 162CA	73.53	25 <b>CB</b>	25C19 23CA	82.52
	25C19 230	74.24	25CB	25C19 23C	70.22
	25C19 162N			25C19 25N	30.03
	25C19 19NE2			25C19 163N	
				25C19 162CB	72.32
162CA	25C19 161C	31.82	162CA	25C19 162N	16.65

			T.	ABLE XI			
162CA	25C19		29.20	162CA	25C19	162CE1	53.97
162CA	25C19	162CB	17.00	161C	25C19	162N	15.17
161C	25C19	163N	54.26	161C	25C19	162CE1	84.02
161C	25C19	162CB	41.75	23CA	25C19	230	29.88
23CA	25C19	23C	18.55	23CA	25C19	25N	55.13
23CA	25C19	19NE2	45.62	230	25C19	23C	14.86
230	25C19	25N	44.51	230	25C19	19NE2	58.93
23C	25C19	2 <b>5N</b>	40.60	23C	25C19	19NE2	45.52
162N	25C19	163N	41.46	162N	25C19	162CE1	69.54
162N	25C19	162CB	28.25	25N	25C19	19NE2	36.78
25N	25C19	163N	74.96	25N	25C19	162CE1	67.79
19NE2	25C19	163N	96.25	19NE2	25C19	162CE1	59.92
163N	25C19	162CE1	51.59	163N	25C19	162CB	39.57
162CE1	25C19	162CB	42.53	1610	25C20	25SG	71.78
1610	25C20	161C	1.07	25 <b>SG</b>	25C20	23CA	80.86
25SG	25C20	230	69.40	25SG	25C20	23C	68.41
25 <i>S</i> G	25C20	161C	71.72	23CA	25C20	230	30.94
23CA	25C20	23C	18.34	230	25C20	23C	14.77
1610	25C21	161C	6.69	1610	25C22	161C	13.34
1610	25C22	161CA	34.90	1610	25C22	161CB	44.27
161C	25C22	161CA	21.77	161C	25C22	161CB	34.39
161CA	25C22	161CB	20.42	184NE1	25 <b>N24</b>	184CZ2	42.68
184NE1	25N24	184CE2	21.38	184NE1	25N24	190 <b>E</b> 1	49.07
184NE1	25N24	162ND1	68.83	184NE1	25 <b>N24</b>	162CE1	51.55
184NE1	25N24	19CD	58.30	184NE1	25 <b>N24</b>	19NE2	75.34
184NE1	25N24	184CD1	9.07	184NE1	25N24	184CH2	51.38
184CZ2	25N24	184CE2	21.94	184CZ2	25N24	190E1	85.09
184CZ2	25N24	162ND1	64.71	184CZ2	25N24	162CE1	57.67
184CZ2	25N24	19CD	98.31	184CZ2	25 <b>N24</b>	184CD1	50.33
184CZ2	25N24	184CH2	9.57	184CE2	25 <b>N24</b>	190E1	68.35
184CE2	25N24	162ND1	68.77	184CE2	25 <b>N24</b>	162CE1	55.33
184CE2	25 <b>N24</b>	19CD	79.25	184CE2	25N24	19NE2	96.20
184CE2	25N24	184CD1	28.46	184CE2	25N24	184CH2	30.12
190E1		162ND1	58.04	190E1	25 <b>N24</b>	162CE1	44.62
190E1	25N24	19CD	16.03	190E1	25 <b>N24</b>	19NE2	29.68
190E1		184CD1	47.21	190E1		184CH2	94.65
162ND1		162CE1	18.14	162ND1	25N24	19CD	72.35
162ND1	25N24	19NE2	76.87	162ND1	25 <b>N24</b>	184CD1	75.84
162ND1	25N24	184CH2	69.90	162CE1	25N24	19CD	60.33
162CE1	25N24	19NE2	69.25	162CE1	25N24	184CD1	58.03
162CE1		184CH2	65.20	19CD	25N24	19NE2	17.04
19CD	25N24	184CD1	53.84	19NE2		184CD1	70.73
184CD1	25N24	184CH2	58.50	25SG	25C25	25CB	32.75
25SG	25C25	25N	68.12	25 <b>S</b> G	25C25	25CA	46.11

			TA	BLE XI			
25 <b>S</b> G	25C25	19NE2	91.26	25\$G	25C25	162ND1	50.45
25SG	25C25	1610	83.98	25SG	25C25	26N	50.26
25SG	25C25	25C	39.24	25 <i>S</i> G	25C25	24C	75.91
25SG	25C25	163N	26.06	25 <i>S</i> G	25C25	190E1	68.77
25SG	25C25	162CA	52.77	25SG	25C25	162CE1	48.87
25SG	25C25	24CA	93.64	25SG	25C25	19CD	79.92
25CB	25C25	25N	43.75	25CB	25C25	25CA	22.80
25CB	25C25	23C	96.85	25CB	25C25	19NE2	58.71
25CB	25C25	162ND1	52.80	25CB	25C25	24N	79.85
25CB	25C25	26N	49.57	25CB	25C25	25C	32.45
25CB	25C25	24C	53.95	25CB	25C25	163N	56.86
25CB	25C25	190E1	39.72	25CB	25C25	162CA	76.70
25CB	25C25	162CE1	41.55	25CB	25C25	24CA	71.02
25CB `	25C25	19CD	48.30	25N	25C25	230	61.20
25N	25C25	25CA	22.53	25N	25C25	23C	54.87
25N	25C25	23CA	71.33	25N	25C25	19NE2	46.58
25N	25C25	162ND1	94.33	25N	25C25	24N	38.82
25N	25C25	26N	37.62	25N	25C25	25C	32.96
25N	25C25	24C	10.58	25N	25C25	163N	94.01
25N	25C25	190E1	53.89	25N	25C25	162CE1	80.69
25N	25C25	24CA	27.34	25N	25C25	19CD	48.28
230	25C25	25CA	82.78	230	25C25	23C	18.78
230	25C25	23CA	35.84	230	25C25	19NE2	73.70
230	25C25	24N	31.06	230	25C25	26N	72.39
230	25C25	25C	83.36	230	25C25	24C	51.02
230	25C25	190E1	99.77	230	25C25	24CA	34.14
230	25C25	19CD	86.19	25CA	25C25	23C	77.38
25CA	25C25	23CA	92.63	25CA	25C25	19NE2	55.22
25CA	25C25	162ND1	75.50	25CA	25C25	24N	61.07
25CA	25C25	26N	33.30	25CA	25C25	25C	18.88
25CA	25C25	24C	31.91	25CA	25C25	163N	72.17
25CA	25C25	190E1	48.70	25CA	25C25	162CA	96.87
25CA	`25C25	162CE1	63.50	25CA	25C25	24CA	49.61
25CA	25C25	19CD	50.26	23C	25C25	23CA	22.22
23C	25C25	19NE2	55.76	23C	25C25	24N	17.04
23C	25C25	26N	77.83	23C	25C25	25C	83.98
23C	25C25	24C	46.58	23C	25C25		82.85
23C	25C25	24CA	28.93	23C	25C25		68.78
23CA	25C25	19NE2	53.36	23CA	25C25	24N	33.27
23CA	25C25	26N	99.45	23CA	25C25	24C	65.16
23CA	25C25	190E1	80.77	23CA	25C25	24CA	48.92
23CA	25C25	19CD	66.76	19NE2	25C25	162ND1	78.91
19NE2	25C25	24N	43.72	19NE2	25C25	26N	83.54
19NE2	25C25	25C	73.61	19NE2	25C25	24C	50.86

			T.	ABLE XI			
19NE2	25C25	190E1	27.88	19NE2	25C25	162CE1	65.60
19NE2	25C25	24CA	51.46	19NE2	25C25	19CD	13.57
162ND1	25C25	1610	66.11	162ND1	25C25	26N	97.49
162ND1	25C25	25C	81.90	162ND1	25C25	163N	47.00
162ND1	25C25	190E1	52.01	162ND1	25C25	162CA	38.64
162ND1	25C25	162CE1	14.60	162ND1	25C25	19CD	65.91
1610	25C25	163N	58.15	1610	25C25	162CA	33.71
1610	25C25	162CE1	80.71	24N	25C25	26N	67.35
24N	25C25	25C	70.00	24N	25C25	24C	31.89
24N	25C25	190E1	68.76	24N	25C25	24CA	16.51
24N	25C25	19CD	55.45	26N	25C25	25C	17.11
26N	25C25	24C	37.71	26N	25C25	163N	70.13
26N	25C25	190E1	81.99	26N	25C25	162CA	99.71
26N	25C25	162CE1	89.75	26N	25C25	24CA	50.94
26N	25C25	19CD	82.06	25C	25C25	24C	38.11
25C	25C25	163N	63.33	25C	25C25	190E1	66.63
25C	25C25	162CA	91.85	25C	25C25	162CE1	73.03
25C	25C25	24CA	55.12	25C	25C25	19CD	69.13
24C	25C25	190E1	62.72	24C	25C25	162CE1	91.22
24C	25C25	24CA	17.90	24C	25C25	19CD	55.23
163N	25C25	190E1	86.02	163N	25C25	162CA	29.58
163N	25C25	162CE1	53.83	163N	25C25	19CD	99.46
190E1	25C25	162CA	90.22	190E1	25C25	162CE1	38.06
190E1	25C25	24CA	71.30	190E1	25C25	19CD	14.33
162CA	25C25	162CE1	52.16	162CE1	25C25	19CD	52.25
24CA	25C25	19CD	60.28	25SG	25026	25N	75.08
25SG	25026	25CB	37.30	25 <b>S</b> G	25026	25CA	54.02
25SG	25026	24C	86.45	25SG	25026	19CD	97.95
25\$G	25026	190E1	82.42	25 <b>S</b> G	25026	25C	45.95
25SG	25026	26N	51.16	25SG	25026	162ND1	46.91
25SG	25026	162CE1	52.51	25N	25026	23C	71.27
25N	25026	25CB	49.91	25N	25026	23CA	95.71
25N	25026	19NE2	63.43	25N	25026	230	74.28
25N	25026	24N	51.65	25N	25026	25CA	24.00
25N	25026	24C	13.92	25N	25026	19CD	63.32
25N	25026	190E1	66.78	25N	25026	24CA	35.47
25N	25026	23N	92.49	25N	25026	220	75.23
25N	25026	25C	29.57	25N	25026	26N	34.00
25N		162ND1	97.43	25N	25026	22C	83.38
25N	25026	162CE1	86.18	23C	25026	23CA	29.24
23C	25026	19NE2	77.74	23C	25026	230 2502	22.63
23C	25026	24N	23.28	23C	25026	25CA	95.23
23C	25026	24C	57.44	23C	25026	19CD	92.50
23C	25026	24CA	35.81	23C	25026	23N	31.40

TABLE XI											
23C	25026	220	45.44	23C	25026	25C	94.92				
23C	25026	26N	85.14	23C	25026	22C	36.80				
25CB	25026	19NE2	75.38	25CB	25026	25CA	26.07				
25CB	25026	24C	63.79	25CB	25026	19CD	62.61				
25CB	25026	190E1	50.03	25CB	25026	24CA	85.37				
25CB	25026	25C	32.00	25CB	25026	26N	48.03				
25CB	25026	162ND1	48.42	25CB	25026	162CE1	40.75				
23CA	25026	19 <b>NE</b> 2	74.13	23CA	25026	230	44.84				
23CA	25026	24N	44.10	23CA	25026	24C	82.84				
23CA	25026	19CD	89.48	23CA	25026	24CA	61.85				
23CA	25026	23N	7.91	23CA	25026	220	37.03				
23CA	25026	22C	22.99	19NE2	25026	230	99.02				
19NE2	25026	24N	59.57	19NE2	25026	25CA	69.12				
19NE2	25026	24C	64.61	19NE2	25026	19CD	15.61				
19NE2	25026	190E1	33.20	19NE2	25026	24CA	66.27				
19NE2	25026	23N	66.26	19NE2	25026	220	37.24				
19NE2	25026	25C	85.04	19NE2	25026	26N	96.39				
19NE2	25026	162ND1	87.10	19NE2	25026	22C	51.16				
19NE2	25026	162CE1	72.58	230	25026	24N	40.19				
230	25026	25CA	95.80	230	25026	24C	61.05				
230	25026	24CA	42.72	230	25026	23N	49.93				
230	25026	220	68.06	230	25026	25C	89.22				
230	25026	26N	75.10	230	25026	22C	58.63				
24N	25026	25CA	75.32	24N	25026	24C	38.83				
24N	25026	19CD	72.68	24N	25026	190E1	88.39				
24N	25026	24CA	19.05	24N	25026	23N	41.71				
24N	25026	220	37.44	24N	25026	25C	79.23				
24N	25026	26 <b>N</b>	74.55	24N	25026	22C	37.15				
25CA	25026	24C	37.79	25CA	25026	19CD	62.07				
25CA	25026	190E1	57.50	25CA	25026	24CA	59.43				
25CA	25026	220	92.92	25CA	25026	25C	16.48				
25CA	25026	26N	31.94	25CA	25026	162ND1	74.28				
25CA	25026	162CE1	64.77	24C	25026	19CD	68.53				
24C	25026	190E1	75.73	24C	25026	24CA	21.65				
24C	25026	23N	80.40	24C	25026	220	67.31				
24C	25026	25C	40.54	24C	25026	26N	39.44				
24C	25026	22C	73.15	24C	25026	162CE1	99.55				
19CD	25026	190E1	17.61	19CD	25026	24CA	75.72				
19CD	25026	23N	81.58	19CD	25026	220	52.77				
19CD	25026	25C	78.56	19CD	25026	26N	92.72				
19CD	25026	162ND1	71.98	19CD	25026	22C	66.59				
19CD	25026	162CE1	57.18	1.90E1	25026	24CA	87.94				
190E1	25026	23N	98.65	190E1	25026	220	70.22				
190E1	25026	25C	73.17	190E1	25026	26N	89.43				

TABLE XI												
190E1	25026	162ND1	54.76	190E1	25026	22C	83.88					
190E1	25026	162CE1	39.72	24CA	25026	23N	60.38					
24CA	25026	220	54.26	24CA	25026	25C	60.91					
24CA	25026	26N	55.50	24CA	25026	22C	56.01					
23N	25026	220	29.27	23N	25026	22C	15.20					
220	25026	22C	14.07	25C	25026	26N	16.90					
25C	25026	162ND1	78.47	25C	25026	162CE1	72.68					
26N	25026	162ND1	91.53	26N	25026	162CE1	88.05					
162ND1	25026	162CE1	15.41	25SG	25C27	25N	57.08					
25 <i>S</i> G	25C27	25CB	21.02	25SG	25C27	26N	58.38					
25SG	25C27	25CA	39.38	25SG	25C27	24N	91.00					
25SG	25C27	24C	70.32	25SG	25C27	25C	43.75					
25SG	25C27	1610	68.95	25SG	25C27	26CB	82.20					
25 <b>S</b> G	25C27	24CA	87.49	25SG	25C27	26CG	96.26					
230	25 <b>C27</b>	23C	18.46	230	25C27	25N	62.85					
230	25C27	65CA	58.88	230	25C27	25CB	97.75					
230	25C27	26CD1	56.83	230	25C27	26N	82.95					
230	25C27	23CA	34.44	230	25C27	25CA	81.88					
230	25C27	24N	28.16	230	25C27	24C	51.85					
230	25C27	25C	88.07	230	25C27	65N	46.70					
230	25C27	26CB	86.79	230	25C27	6 <b>6N</b>	79.07					
230	25C27	24CA	33.37	230	25C27	26CG	69.96					
230	25C27	65C	66.97	23C	25C27	25N	53.34					
23C	25C27	65CA	76.45	23C	25C27	25CB	84.25					
23C	25C27	26CD1	70.76	23C	25C27	26N	83.38					
23C	25C27	23CA	21.08	23C	25C27	25CA	72.15					
23C	25C27	24N	15.52	23C	25C27	24C	46.42					
23C	25C27	25C	83.42	23C	25C27	65N	62.90					
23C	25C27	26CB	96.45	23C	25C27	66N	97.46					
23C	25C27	24CA	29.98	23C	25C27	26CG	81.94					
23C	25C27	65C	85.36	25 <b>N</b>	25C27	25CB	36.48					
25N	25C27	26CD1	68.71	25 <b>N</b>	25C27	26N	39.49					
25N	25C27	23CA	66.25	25 <b>N</b>	25C27	25CA	19.15					
25N	25C27	24N	37.84	25N	25C27	24C	14.42					
25N	25C27	25C	31.97	25N	25C27	26CB	68.32					
25N	25C27	24CA	30.42	25N	25C27	26CG	67.30					
65CA	25C27	26CD1	53.12	65CA	25C27	23CA	82.66					
65CA	25C27	24N	86.51	65CA	25C27	24C	98.45					
65CA	25C27	65N	15.90	65CA	25C27	26CB	77.82					
65CA	25C27	66N	30.25	65CA	25C27	24CA	84.16					
65CA	25C27	26CG	63.87	65CA	25C27	65C	14.85					
25CB	25C27	26CD1	96.07	25CB	25C27	26N	48.98					
25CB	25C27	23CA	83.62 70.10	25CB	25C27	25CA	20.65					
25CB	25C27	24N	70.19	25CB	25C27	24C	50.31					

			т	ABLE XI			
25CB	25C27	25C	32.85	25CB	25C27	1610	88.13
25CB	25C27	26CB	78.77	25CB	25C27	24CA	66.76
25CB	25C27	26CG	88.03	26CD1	25C27	26N	49.41
26CD1	25C27	23CA	90.70	26CD1	25C27	25CA	76.35
26CD1	25C27	24N	67.23	26CD1	25C27	24C	55.82
26CD1	25C27	25C	64.70	26CD1	25C27	65N	58.91
26CD1	25C27	26CB	32.69	26CD1	25C27	66N	42.33
26CD1	25C27	24CA	52.50	26CD1	25C27	26CG	14.60
26CD1	25C27	65C	44.57	26N	25C27	25CA	33.28
26N	25C27	24N	69.36	26N	25C27	24C	38.31
26N	25C27	25C	16.13	26N	25C27	26CB	31.48
26N	25C27	6 <b>6N</b>	83.54	26N	25C27	24CA	53.43
26N	25C27	26CG	39.23	26N	25C27	65C	91.88
23CA	25C27	25CA	82.90	23CA	25C27	24N	32.54
23CA	25C27	24C	63.47	23CA	25C27	25C	98.17
23CA	25C27	65N	67.00	23CA	25C27	24CA	49.60
23CA	25C27	65C	95.04	25CA	25C27	24N	56.77
25CA	25C27	24C	31.07	25CA	25C27	25C	19.16
25CA	25C27	26CB	64.76	25CA	25C27	24CA	48.93
25CA	25C27	26CG	70.17	24N	25C27	24C	31.48
24N	25C27	25C	68.08	24N	25C27	65N	74.86
24N	25C27	26CB	87.15	24N	25C27	24CA	17.56
24N	25C27	26CG	75.54	24N	25C27	65C	92.54
24C	25C27	25C	37.23	24C	25C27	65N	93.14
24C	25C27	26CB	61.81	24C	25C27	66N	98.06
24C	25C27	24CA	18.48	24C	25C27	26CG	56.79
24C	25C27	65C	96.78	25C	25C27	26CB	46.70
25C	25C27	6 <b>6N</b>	99.53	25C	25C27	24CA	55.28
25C	25C27	26CG	<b>55.</b> 30	65N	25C27	26CB	88.24
65N	25C27	66N	45.67	6 <b>5N</b>	25C27	24CA	76.36
65N	25C27	26CG	72.00	6 <b>5N</b>	25C27	65C	30.06
26CB	25C27	6 <b>6N</b>	53.44	26CB	25C27	24CA	69.63
26CB	25C27	26CG	18.09	26CB	25C27	65C	64.59
66N		24CA	91.56		25C27		45.61
66N	25C27	65C	15.61	24CA	25C27		58.92
24CA	25C27	65C	85.96	2 SCG	25C27		52.65
25SG		1610	76.60	25 <b>S</b> G	25028		80.25
25 <i>S</i> G	25028	26CD1	87.41	25SG	25028		47.59
25SG	25028	26CB	75.36	25 <i>S</i> G	25028		77.06
25 <i>S</i> G	25028	25CB	9.94	25SG	25028		39.01
65CA	25028	230	50.13	65CA	25028	6611	33.41
65CA	25028	660	67.54				
65CA	25028	65C		65CA		_	
65CA	25C28	26CB	76.12	1610	25028	161C	10.43

			T	ABLE XI			
1610	25028	25CB	85.93	1610	25028	163N	57.90
230	25028	66N	73.33	230	25028	660	98.34
230	25028	26CD1	46.36	230	25028	65C	62.14
230	25028	26N	63.88	230	25028	26CB	73.66
230	25028	25CB	70.99	6 <u>6</u> N	25028	660	34.18
6 <b>6</b> N	25028	26CD1	42.77	66N	25028	65C	16.09
66N	25028	26N	80.82	66 <b>N</b>	25028	26CB	55.05
660	25028	26CD1	53.93	660	25028	65C	50.27
660	25028	26N	71.20	660	25028	26CB	41.80
660	25028	163N	96.32	26CD1	25028	65C	44.16
26CD1	25028	26N	42.70	26CD1	25028	26CB	31.40
26CD1	25028	25CB	78.43	65C	25028	26N	86.36
65C	25028	26CB	65.21	26N	25028	26CB	29.67
26N	25028	25CB	40.69	26N	25028	163N	65.13
26CB	25028	25CB	69.78	26CB	25028	163N	79.13
161C	25028	25CB	86.86	161C	25028	163N	52.19
25CB	25028	163N	47.47	660	25C29	6 <b>6N</b>	48.91
660	25C29	65CA	91.41	660	25C29	65C	68.29
660	25C29	66C	13.27	660	25C29	26CD1	67.50
660	25C29	66CA	34.11	660	25C29	26CB	50.71
660	25C29	26CG	54.55	660	25C29	65N	99.08
660	25C29	650	66.95	660	25C29	26N	80.09
66N	25C29	65CA	42.87	66N	25C29	65C	19.38
66N	25C29	66C	36.24	66N	25C29	26CD1	51.44
66N	25C29	66CA	15.57	66N	25C29	26CB	68.44
6 <b>6N</b>	25C29	26CG	54.31	6 <b>6N</b>	25C29	230	78.55
6 <b>6N</b>	25C29	65N	50.66	6 <b>6N</b>	25C29	650	18.11
66N	25C29	26N	91.42	65CA	25C29	65C	23.73
65CA	25C29	66C	79.07	65CA	25C29	26CD1	55.65
65CA	25C29	66CA	58.35	65CA	25C29	26CB	89.86
65CA	25C29	26CG	70.46	65CA	25C29	230	47.25
65CA	25C29	65N	7.81	65CA	25C29	650	25.70
65CA .	25C29	26N	94.75	65C	25C29	66C	55.59
65C	25C29	26CD1	52.52	65C	25C29	66CA	34.70
65C	25C29	26CB	79.47	65C	25C29	26CG	61.82
65C	25C29	230	65.09	65C	25C29	65N	31.53
65C	25C29	650	2.78	65C	25C29	26N	95.49
66C	25C29	26CD1	64.26	66C	25C29	66CA	21.04
66C	25C29	26CB	56.28	66C	25C29	26CG	54.75
66C	25C29	65N	36.84	66C	25C29	650	54.09
66C	25C29	26N	86.08	26CD1	25C29	66CA	57.66
26CD1	25C29	26CB	35.62	26CD1	25C29	25SG	81.23
26CD1	25C29	26CG	17.23	26CD1	25C29	230	45.73
26CD1	25C29	65N	58.39	26CD1	25C29	650	54.62

			T	ABLE XI			
26CD1	25C29	26N	42.97	66CA	25C29	26CB	64.36
66CA	25C29	26CG	54.97	66CA	25C29	230	92.04
66CA	25C29	65N	66.15	66CA	25C29	650	33.08
66CA	25C29	26N	91.66	26CB	25C29	25 <b>S</b> G	73.09
26CB	25C29	26CG	19.41	26CB	25C29	230	74.83
26CB	25C29	65N	93.60	26CB	25C29	650	80.55
26CB	25C29	26N	29.84	25 <b>S</b> G	25C29	26CG	81.03
25SG	25C29	230	65.42	25 <b>S</b> G	25C29	1610	57.72
25SG	25C29	26N	44.40	26CG	25C29	230	61.35
26CG	25C29	65N	74.27	26CG	25C29	650	63.28
26CG	25C29	26N	37.32	230	25C29	65N	42.49
230	25C29	650	67.82	230	25C29	26N	59.60
65N	25C29	650	33.46	65N	25C29	26N	94.60
650	25C29	26N	97.54	660	25C30	66N	43.16
660	25C30	66C	13.69	660	25C30	65CA	74.47
660	25C30	66CA	31.15	660	25C30	65C	58.33
66N	25C30	66C	35.28	6 <b>6N</b>	25C30	65CA	33.44
66N	25C30	66CA	17.60	6 <b>6N</b>	25C30	65C	15.22
66C	25C30	65CA	68.55	66C	25C30	66CA	19.80
66C	25C30	65C	50.23	65CA	25C30	66CA	50.69
65CA	25C30	65C	19.75	66CA	25C30	65C	31.38
660	25C31	66C	9.16	660	25C31	66N	30.51
660	25C31	163CB	81.88	1610	25C31	161C	15.04
1610	25C31	163CB	81.82	1610	25C31	1600	59.31
66C	25C31	66N	28.88	66C	25C31	163CB	90.84
161C	25C31	163CB	80.88	161C	25C31	1600	48.22
6 <b>6N</b>	25C31	163CB	97.56				

TABLE XII

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
184CB	25C1	184CG	21.22	184CB	25C1	180D1	72.94
184CB	25C1	184CD1	35.50	184CB	25C1	184CA	19.59
184CB	25C1	184CD2	33.67	184CB	25C1	1840	34.28
184CB	25C1	184C	30.67	184CG	25C1	180D1	81.92
184CG	25C1	184CD1	18.31	184CG	25C1	184CA	34.17
184CG	25C1	184CD2	16.99	184CG	25C1	1840	55.47
184CG	25C1	184C	50.07	180D1	25C1	184CD1	74.79
180D1	25C1	184CA	53.36	180D1	25C1	184CD2	98.79
180D1	25C1	1840	61.42	180D1	25C1	184C	50.27
184CD1	25C1	184CA	39.83	184CD1	25C1	184CD2	28.30
184CD1	25C1	1840	67.35	184CD1	25C1	184C	57.99
184CA	25C1	184CD2	50.19	184CA	25C1	1840	29.82
184CA	25C1	184C	18.22	184CD2	25C1	1840	66.46
184CD2	25C1	184C	64.29	1840	25C1	184C	14.58
180D1	25C2	184CD1	93.56	180D1	25C2	184CB	85.20
180D1	25C2	184CA	63.93	180D1	25C2	184CG	97.27
180D1	25C2	18CG	7.95	180D1	25C2	200	82.27
180D1	25C2	20 <b>N</b>	45.49	180D1	25C2	184C	56.77
180D1	25C2	18ND2	22.89	180D1	25C2	20CA	55.50
180D1	25C2	1840	66.85	180D1	25C2	20C	72.94
180D1	25C2	19CG	66.66	184CD1	25C2	184CB	36.96
184CD1	25C2	184CA	43.35	184CD1	25C2	184CG	19.17
184CD1	25C2	200	93.58	184CD1	25C2	20N ·	91.27
184CD1	25C2	184C	62.15	184CD1	25C2	184NE1	14.42
184CD1	25C2	1840	68.85	184CD1	25C2	19CG	47.81
184CB	25C2	184CA	21.51	184CB	25C2	184CG	20.93
184CB	25C2	18CG	90.69	1.84CB	25C2	184C	32.58
184CB	25C2	184NE1	48.17	184CB	25C2	1840	33.58
184CB	25C2	1.9CG	75.81	184CA	25C2	184CG	35.93
184CA	25C2	18CG	69.84	184CA	25C2	20N	91.24
184CA	25C2	184C	19.09	184CA	25C2	18ND2	85.37
184CA	25C2	184NE1	57.50	184CA	25C2	1840	29.61
184CA	25C2	19CG	65.69	184CG	25C2	184C	51.91
184CG	25C2	184NE1	27.66	184CG	25C2	1840	54.51
184CG	25C2	19CG	65.63	18CG	25C2	200	83.37
18CG	25C2	20N	47.67	18CG	25C2	184C	60.57

			т	ABLE XII			
18CG	25C2	18ND2	15.73	18CG	25C2	20CA	54.96
18CG	25C2	1840	69.03	18CG	25C2	20C	72.82
18CG	25C2	19CG	73.59	200	25C2	20N	37.03
200	25C2	18ND2	77.52	200	25C2	20CA	29.71
200	25C2	184NE1	87.40	200	25C2	20C	13.14
200	25C2	19CG	52.69	20N	25C2	184C	94.70
20N	25C2	18ND2	46.43	20N	25C2	20CA	17.58
20N	25C2	184NE1	94.15	20 <b>N</b>	25C2	20C	29.99
20N	25C2	19CG	44.09	184C	25C2	18ND2	74.66
184C	25C2	184NE1	76.04	184C	25C2	1840	14.64
184C	25C2	19CG	80.33	18ND2	25C2	20CA	47.82
18ND2	25C2	1840	81.12	18ND2	25C2	20C	65.32
18ND2	25C2	19CG	81.86	20CA	25C2	20C	17.90
20CA	25C2	19CG	58.30	184NE1	25C2	1840	81.39
184NE1	25C2	20C	98.78	184NE1	25C2	19CG	50.29
1840	25C2	19CG	94.21	20C	25C2	19CG	58.17
200	25C3	180D1	94.92	200	25C3	20C	14.93
200	25C3	20N	45.22	200	25C3	20CA	34.91
200	25C3	19CG	67.31	200	25C3	18CG	90.10
200	25C3	19C	44.11	200	25C3	19CD	68.85
180D1	25C3	20C	83.09	180D1	25C3	20N	50.73
180D1	25C3	184CD1	86.70	180D1	25C3	20CA	61.59
180D1	25C3	19CG	73.40	180D1	25C3	184CG	83.83
180D1	25C3	18CG	7.01	180D1	25C3	184CB	69.00
180D1	25C3	184CA	52.51	180D1	25C3	19C	56.82
180D1	25C3	19CD	89.80	20C	25C3	20N	36.79
20C	25C3	20CA	21.55	20C	25C3	19CG	72.42
20C	25C3	18CG	77.56	20C	25C3	19C	39.99
20C	25C3	19CD	77.92	20N	25C3	20CA	20.57
20N	25C3	19CG	52.06	20N	25C3	18CG	47.19
20N	25C3	184CA	88.94	20N	25C3	19C	12.18
20N	25C3	19CD	65.92	184CD1	25C3	19CG	54.41
184CD1	25C3	184CG	17.14	184CD1	25C3	184NE1	17.22
184CD1		18CG			25C3		32.16
184CD1	25C3	184CA	38.24	184CD1	25C3	19C	92.25
184CD1	25C3	19CD	49.63 56.01	20CA	25C3		69.55
20CA 20CA	25C3	18CG 19CD		20CA	25C3		29.66 69.80
20CA 19CG	25C3 25C3	19CD 184NE1	80.70 57.02	19CG	25C3	184CG 18CG	
			76.30	19CG	25C3		76.04
19CG 19CG	25C3	184CB 19C	40.46	1.9CG	25C3	184CA 19CD	66.22 16.92
19CG 184CG	25C3 25C3	19C 184NE1	28.96	19CG 184CG	25C3 25C3		90.83
184CG	25C3	184CB	18.06	184CG	25C3	184CA	31.35
184CG	25C3	19CD	66.65	184ME1	25C3	184CB	46.68
T0-4012	2343	1000	55.55	エウオロセナ	2, 3 (-)	TO:#CD	-U.00

			T	ABLE XII			
184NE1	25C3	184CA	55.31	184NE1	25C3	19C	97.43
184NE1	25C3	19CD	46.53	18CG	25C3	184CB	75.84
18CG	25C3	184CA	59.51	18CG	25C3	19C	54.70
18CG	25C3	19CD	92.81	184CB	25C3	184CA	18.20
184CB	25C3	19CD	77.96	184CA	25C3	19C	86.12
184CA	25C3	19CD	72.78	19C	25C3	19CD	53.79
200	25C4	20C	9.86	200	25C4	19CG	57.67
200	25C4	180D1	69.35	200	25C4	20N	32.43
184CD1	25C4	184NE1	18.35	184CD1	25C4	184CG	15.96
184CD1	25C4	19CG	48.28	184CD1	25C4	184CE2	26.62
184CD1	25C4	180D1	68.05	184CD1	25C4	20N	82.80
184NE1	25C4	184CG	27.99	184NE1	25C4	19CG	53.65
184NE1	25C4	184CE2	15.20	184NE1	25C4	180D1	85.79
184NE1	25C4	20N	94.26	20C	25C4	19CG	62.70
20C	25C4	180D1	63.89	20C	25C4	20N	29.77
184CG	25C4	19CG	62.53	184CG	25C4	184CE2	27.11
184CG	25C4	180D1	67.71	184CG	25 <b>C4</b>	20N	91.33
19CG	25C4	184CE2	68.70	19CG	25C4	180D1	57.18
19CG	25C4	20N	42.85	184CE2	25C4	180D1	93.39
180D1	25C4	20N	37.93	184CD1	25C5	184NE1	17.38
184CD1	25C5	200	82.23	184CD1	25C5	184CG	16.75
184CD1	25C5	184CE2	27.59	184CD1	25C5	184CD2	27.17
184NE1	25C5	200	86.19	184NE1	25C5	184CG	27.91
184NE1	25C5	184CE2	16.68	184NE1	25C5	184CD2	27.27
200	25C5	184CG	96.36	184CG	25C5	184CE2	27.86
184CG	25C5	184CD2	16.99	184CE2	25C5	184CD2	16.85
184CG	25C6	184CD1	17.30	184CG	25C6	184CD2	18.02
184CG	25C6	184CB	18.64	184CG	25C6	184NE1	27.34
184CG	25C6	184CE2	27.75	184CD1	25C6	184CD2	28.13
184CD1	25C6	184CB	32.10	184CD1	25C6	184NE1	16.25
184CD1	25C6	184CE2	26.86	184CD2	25C6	184CB	32.59
184CD2	25C6	184NE1	27.14	184CD2	25C6	184CE2	16.75
184CB	25C6	184NE1	45.42	184CB	25C6	184CE2	45.85
184NE1	25C6	184CE2	16.05	200	25C7	20C	5.96
200	25C7	21CA	34.09	200	25C7	19CD	73.49
200	25C7	19CG	60.92	200	25C7	210E1	66.16
200	25C7	21N	17.58	200	25 <b>C</b> 7	190E1	87.57
200	25C7	19NE2	69.26	20C	25C7	21.CA	31.76
20C	25C7	19CD	79.31	20C	25C7	19CG	66.28
20C	25C7	210E1	60.89	30C	25C7	21N	14.18
20C	25C7	190E1	93.30	20C	25C7	19NE2	75.21
184NE1	25C7	19CD	47.01	184NE1	25C7	1.9CG	52.85
184NE1	25C7	184CD1	16.44	184NE1	25 <b>C7</b>	190E1	33.68
184NE1	25C7	19ME2	59.90	21CA	25C7	19CD	96.04

			TA	ABLE XII			
21CA	25C7	19CG	89.60	21CA	25C7	210E1	41.39
21CA	25C7	21N	17.76	21CA	25C7	19NE2	85.25
19CD	25C7	19CG	18.31	19CD	25C7	21N	88.09
19CD	25C7	184CD1	47.08	19CD	25C7	190E1	14.53
19CD	25C7	19NE2	15.43	19CG	25C7	21N	77.54
19CG	25C7	184CD1	46.14	19CG	25C7	190E1	28.58
19CG	25C7	19NE2	28.52	210E1	25C7	21N	49.85
21N	25C7	19NE2	80.92	184CD1	25C7	190E1	37.45
184CD1	25C7	19NE2	62.12	190E1	25C7	19NE2	26.24
200	2508	19CD	90.86	200	2508	19NE2	87.53
200	2508	19CG	72.09	200	2508	20C	2.00
200	2508	220	62.72	200	2508	19CB	63.81
19CD	2508	190E1	19.87	19CD	2508	19NE2	21.15
19CD	2508	19CG	23.15	19CD	2508	184NE1	60.87
19CD	2508	184CD1	58.07	19CD	2508	20C	89.41
19CD	2508	220	57.75	19CD	2508	184CE2	73.67
19CD	2508	19CB	27.32	190E1	2508	19NE2	35.82
190E1	2508	19CG	37.30	190E1	2508	184NE1	43.39
190E1	2508	184CD1	45.88	190E1	2508	220	74.89
190E1	2508	184CE2	55.38	190E1	2508	19CB	45.39
19NE2	2508	19CG	37.34	19NE2	2508	184NE1	79.21
19NE2	2508	184CD1	78.91	19NE2	2508	20C	85.68
19NE2	2508	220	39.13	19NE2	2508	184CE2	91.05
19NE2	2508	19CB	33.61	19CG	2508	184NE1	65.75
19CG	2508	184CD1	54.97	19CG	2508	20C	71.02
19CG	2508	220	61.28	19CG	2508	184CE2	78.82
19CG	2508	19CB	12.33	184NE1	2508	184CD1	18.60
184NE1	2508	184CE2	13.25	184NE1	2508	19CB	77.87
184CD1	2508	184CE2	27.58	184CD1	2508	19CB	67.24
20C	2508		60.75	20C	2508	19CB	62.46
220	2508	19CB	49.98	184CE2	2508	19CB	91.01
190E1	25C9	19NE2	33.38	190E1	25C9	19CD	18.07 84.29
190E1	.25C9	184NE1	40.26	190 <b>E1</b>	25C9	200 220	
190E1	25C9	19CG	30.18	190E1	25C9	184CD1	69.67 39.42
190E1	25C9	184CE2	53.38	190E1	25C9	184NE1	73.02
19NE2	25C9	19CD	19.36	19NE2	25C9	19CG	30.40
19NE2	25C9	200	69.77	19NE2	25C9	19CG 184CE2	86.66
19NE2	25C9	220	36.73	19NE2	25C9 25C9	184NE1	54.70
19NE2	25C9	184CD1	68.70	19CD		19CG	17.06
19CD	25C9	200	69.22	19CD	25C9 25C9	19CG 184CE2	68.86
19CD	25C9	220	53.02	19CD		200	95.14
19CD	25C9	184CD1	49.36	184NE1	25C9	184CE2	14.51
184NE1	25C9	19CG	55.54	184ME1	25C9		
184NE1	25C9	184CD1	14.36	200	25C9	19CG	54.13

			T	ABLE XII			•
200	25C9	220	54.06	200	25C9	184CD1	80.80
19CG	25C9	220	53.71	19CG	25C9	184CE2	69.95
19CG	25C9	184CD1	45.63	220	25C9	184CD1	99.26
184CE2	25C9	184CD1	26.16	19NE2	25010	200	62.12
19NE2	25010	220	35.56	19NE2	25010		15.61
19NE2	25010	190E1	26.78	200	25010	220	52.61
200	25010	19CD	59.60	200	25010	190E1	70.63
220	25010	19CD	48.47	220	25010	190E1	61.86
19CD	25010	190E1	14.34	162ND1	25C11	184CZ2	66.42
162ND1	25C11	184NE1	66.34	162ND1	25C11	162CE1	17.78
162ND1	25C11	190E1	57.83	162ND1	25C11	184CE2	67.52
162ND1	25C11	162CG	14.11	162ND1	25C11	19NE2	72.38
162ND1	25C11	19CD	68.15	184CZ2	25C11	184NE1	34.50
184CZ2	25C11	162CE1	56.59	184CZ2	25C11	190E1	69.19
184CZ2	25C11	184CE2	17.54	184CZ2	25C11	162CG	60.04
184CZ2	25C11	19NE2	95.66	184CZ2	25C11	19CD	80.47
184NE1	25C11	162CE1	49.27	184NE1	25C11	190E1	37.54
184NE1	25C11		17.27	184NE1	25C11	162CG	68.99
184NE1	25C11	19NE2	62.26	184NE1	25C11	19CD	46.90
162CE1	25C11	190E1	42.46	162CE1	25C11	184CE2	53.20
162CE1	25C11		26.53	162CE1	25C11	19NE2	62.11
162CE1	25C11	19CD	54.33	190E1	25C11	184CE2	53.84
190E1	25C11	162CG	68.94	190E1	25C11	19NE2	26.94
190E1	25C11	19CD	13.30	184CE2	25C11	162CG	65.59
184CE2	25C11	19NE2	79.38	184CE2	25C11	19CD	64.03
162CG	25C11	19NE2	86.02	162CG	25C11	19CD	80.39
19NE2	25C11	19CD	15.36	184CZ2	25C15	184CH2	16.48
184CZ2	25C15	1430E1	72.66	184CZ2	25C15	1370	75.04
184CH2	25C15	1430E1	58.91	184CH2		1370	61.89
1430E1	25C15	1370	63.98	162ND1	25C16	162CE1	19.73
162ND1	25C16	162CG	16.60	162ND1	25C16	25 <b>S</b> G	47.39
162ND1	25C16	1610	72.64	162ND1	25C16	190E1	62.63
162ND1	25C16	162CB	33.51	162ND1	25C16	162CA	39.24
162ND1		162NE2	20.67	162ND1	25C16	19NE2	81.12
162ND1		184CZ2	64.44	162ND1	25C16	25CB	44.50
162ND1		184NE1	64.44	162ND1		162CD2	17.81
162CE1		162CG	32.31	162CE1	25C16	25 <b>S</b> G	53.22
162CE1	25C16		91.93	162CE1	25C16	190El	44.18
162CE1		162CB	51.51	162CE1		162CA	58.97
162CE1		162NE2	11.09	162CE1	25C16	19NE2	66.33
162CE1		184CZ2	53.67	162CE1	25C16	25CB	41.27
162CE1		184NE1	46.11	162CE1		162CD2	24.34
162CG	25C16	25SG	59.98	162CG	25C16		66.72
162CG	25C16	190E1	76.48	162CG	25C16	162CB	19.50

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			TA	ABLE XII			
162CG	25C16	162CA	32.20	162CG	25C16	162NE2	27.13
162CG	25C16	19NE2	97.24	162CG	25C16	184CZ2	60.98
162CG	25C16	25CB	60.57	162CG	25C16	184NE1	70.16
162CG	25C16	162CD2	12.90	25SG	25C16	1610	62.69
25SG	25C16	190E1	65.19	25 <i>S</i> G	25C16	162CB	65.02
25SG	25C16	162CA	52.80	25SG	25C16	162NE2	62.49
25SG	25C16	19NE2	61.43	25 <i>S</i> G	25C16	25CB	20.75
25SG	25C16	184NE1	91.34	25SG	25C16	162CD2	65.19
1610	25C16	162CB	50.15	1610	25C16	162CA	34.64
1610	25C16	162NE2	92.16	1610	25C16	25CB	82.66
1610	25C16	162CD2	79.61	190E1	25C16	162CB	95.63
190E1	25C16	162CA	99.99	190E1	25C16	162NE2	51.09
190E1	25C16		27.38	190E1	25C16	184CZ2	62.46
190E1	25C16		44.47	190E1	25C16	184NE1	33.86
190E1	25C16	162CD2	67.06	162CB	25C16	162CA	18.75
162CB	25C16	162NE2	46.53	162CB	25C16	184CZ2	73.28
162CB	25C16	25CB	72.45	162CB	25C16	184NE1	88.17
162CB	25C16	162CD2	31.28	162CA	25C16	162NE2	57.73
162CA	25C16	184CZ2	91.31	162CA	25C16	25CB	65.83
162CA	25C16		45.08	162NE2	25C16	19NE2	75.48
162NE2	25C16	184CZ2	45.32	162NE2	25C16	25CB	52.16
162NE2	25C16	184NE1	44.44	162NE2	25C16	162CD2	16.02
19NE2	25C16	184CZ2	88.09	19NE2	25C16	25CB	44.94
19NE2	25C16	184NE1	58.55	19NE2	25C16	162CD2	90.64
184CZ2	25C16	25CB	91.81	184CZ2	25C16	184NE1	29.55
184CZ2	25C16	162CD2	49.38	25CB	25C16	184NE1	71.82
25CB	25C16	162CD2	60.66	184NE1	25C16	162CD2	57.37
162ND1	25017	162CG	25.38	162ND1	25017	162CB	48.94
162ND1	25017	162CE1	18.45	162ND1	25017	1610	93.58
162ND1	25017	162CA	54.76	162ND1	25017	25SG	48.39
162ND1	25017	161C	89.31	162ND1	25017	162CD2	25.71
162ND1	25017	162N	73.99	162ND1	25017	162NE2	21.11
162ND1	25017	184CZ2	69.27	162ND1	25017	137CB	75.37
162CG		162CB	26.64	162CG		162CE1	38.41
162CG	25017		86.53	162CG	25017		42.50
162CG	25017	25 <b>S</b> G	67.49	162CG	25017		76.51
162CG		162CD2	13.63	162CG	25017		58.79
162CG		162NE2	29.02	162CG		1610D1	79.16
162CG		184CZ2	67.73	162CG		137CB	51.32
162CB		162CE1	64.60	162CE	25017		65.36
162CB			24.38	162CB	25017	25SG	76.49
162CB	25017	161C	52.48	162CB	25017		38.18
162CB	25017		34.58	162CB		162NE2	55.49
162CB	25017	1610D1	54.52	1.62CB	25017	184CZ2	85.85

			T/	ABLE XII			
162CB	25017	7 137CB	42.56	162CE1	25017	162CA	73.15
162CE1	25017	7 25SG	55.22	162CE1	25017	162CD2	31.50
162CE1	25017	7 162N	92.28	162CE1	25017	162NE2	14.96
162CE1	25017	184CZ2	56.06	162CE1	25017	137CB	81.03
1610	25017	162CA	44.07	1610	25017	25SG	69.63
1610	25017	161C	15.96	1610	25017	162N	31.46
1610	25017	1610D1	56.32	1610	25017	137CB	91.92
162CA	25017	25 <b>S</b> G	61.24	162CA	25017	161C	35.13
162CA	25017	162CD2	56.09	162CA	25017	162N	19.37
162CA	25017	162NE2	69.58	162CA	25017	1610D1	58.28
162CA	25017	137CB	64.45	25SG	25017	161C	77.58
25 <i>S</i> G	25017	162CD2	73.41	25SG	25017	162N	74.60
25SG	25017	162NE2	67.15	161C	25017	162CD2	89.68
161C	25017	162N	17.96	161C	25017	1610D1	42.82
161C	25017	137CB	75.96	162CD2	25017	162N	71.80
162CD2	25017	162NE2	18.45	162CD2	25017	1610D1	86.33
162CD2	25017	184CZ2	54.40	162CD2	25017	137CB	50.73
162N	25017	162NE2	87.31	162N	25017	1610D1	42.68
162N	25017		63.55	162NE2	25017	184CZ2	48.21
162NE2	25017		66.23	1610D1	25017	137CB	44.42
184CZ2	25017		61.47	25SG	25N18	162ND1	50.41
25SG	25N18		53.56	25 <i>S</i> G	25N18	1610	67.92
25SG	25N18		71.90	25 <b>S</b> G	25N18	190E1	69.62
25SG	25N18	25CB	22.82	25SG	25N18	23CA	87.13
25SG	25N18	162CG	56.99	25 <b>S</b> G	25N18	19CD	72.29
162ND1	2 <b>5N18</b>	162CE1	17.36	162ND1	25N18	1610	69.09
162ND1	25 <b>N1</b> 8	19NE2	84.51	162ND1	25N18	190E1	59.48
162ND1	25N18	25CB	49.58	162ND1	25N18	162CG	11.34
162ND1	25N18	19CD	72.69	162CE1	25N18	1610	86.28
162CE1	25N18	19NE2	68.55	162CE1	25 <b>N1</b> 8	190E1	42.23
162CE1	25N18	25CB	43.96	162CE1	25N18	162CG	27.38
162CE1	25N18	19CD	55.80	1610	25N18	25CB	89.34
1610	25N18	162CG	61.14	19 <b>NE</b> 2	25N18	190E1	29.63
19NE2	25N18	25CB	50.72	19NE2	25N18	23CA	55.49
19NE2	25N18	162CG	95.57	19NE2	25N18	19CD	15.35
190E1	25N18	25CB	47.37	19 <b>0E1</b>	25N18	23CA	85.11
190E1		162CG	69.47	190E1	25N18	19CD	14.61
25CB	25N18	23CA	81.33	25CB		162CG	59.74
25CB	25N18	19CD	49.52	23CA	25N18	19CD	70.70
162CG	25N18	19CD	83.18	25 <b>SG</b>	25C19	1610	92.47
25SG	25C19	162ND1	54.14	25SG	25C19	25CB	20.8€
25SG	25C19	162CA	62.74	25 <b>SG</b>	25C19	161C	88.58
25SG	25C19	1.62CE1	51.71	25 <b>S</b> G	25C13	23CA	97.99
25SG	25C19	25N	41.07	25 <b>S</b> G	25C19	23C	82.10

			Т	ABLE XII			
25SG	25C1	9 230	81.30		25C1	9 162CG	59.31
25 <i>S</i> G	25C1		69.11	25SG	25C1	9 1.62CB	67.82
25 <i>S</i> G	25C1		76.57	1610	25C1	9 162ND1	78.79
1610	25C1	9 162CA	38.01	1610	25C1	9 161C	7.13
1610	25C19	9 162CE1	93.42	1610	25C1	9 162CG	66.49
1610	25C19		49.06	1610	25C1		21.75
162ND1	25C19	9 25CB	52.62	162ND1	25C1	9 162CA	42.73
162ND1	25C19	9 161C	71.68	162ND1	25C19	162CE1	14.66
162ND1	25C19	9 25N	83.32	162ND1	25C19	162CG	12.47
162ND1	25C19	9 19NE2	75.83	162ND1	25C19	162CB	30.03
162ND1	25C19	9 162N	57.42	25CB	25C19	162CA	76.45
25CB	25C19	162CE1	44.02	25CB	25C19	23CA	85.20
25CB	25C19		31.12	25CB	25C19	23C	72.96
25CB	25C19		77.28	25CB	25C19	162CG	62.09
25CB	25C19		48.58	25CB	25C19	162CB	75.75
25CB	25C19		92.17	162CA	25C19	161C	31.40
162CA	25C19	162CE1	56.83	162CA	25C19	162CG	31.62
162CA	25C19		17.87	162CA	25C19	162N	16.50
161C	25C19		86.32	161C	25C19	162CG	59.37
161C	25C19	162CB	41.93	161C	25C19	162N	14.94
162CE1	25C19		72.85	162CE1	25C19	162CG	27.09
162CE1	25C19	19NE2	61.17	162CE1	25C19	162CB	44.66
162CE1	25C19		71.95	23CA	25C19	25N	56.99
23CA	25C19		18.49	23CA	25 <b>C</b> 19	230	29.45
23CA	25C19		51.58	25 <b>N</b>	25C19	23C	42.38
25N	25C19		46.35	25N	25C19	162CG	93.20
25N	25C19		42.58	23C	25C19	230	14.74
23C	25C19		52.68	230	25C19	19NE2	66.29
162CG	25C19	19NE2	88.25	162CG	25C19	162CB	17.57
162CG	25C19	162N	45.36	162CB	25C19	162N	28.45
184NE1	25N20	190E1	48.31	184NE1	25 <b>N20</b>	19CD	61.42
184NE1	25N20	184CE2	19.07	184NE1	25N20	19NE2	80.38
184NE1	25N20	184CZ2	38.30	184NE1	25N20	162CE1	53.42
184NE1		184CD1	12.98	184NE1	25 <b>N2</b> 0	162ND1	68.69
184NE1	25N20	19CG	57.21	190E1	25 <b>N2</b> 0	19CD	18.12
190E1	25N20		65.80	190 <b>E1</b>	25N20	19NE2	33.85
190E1		184CZ2	80.73	190E1		162CE1	45.71
190E1		184CD1	44.65	190E1	25N20	162ND1	59.71
190E1	25N20	19CG	26.58	19CD	25N20	184CE2	80.19
19CD	25N20	19NE2	19.18	19CD	25N20	184CZ2	97.13
19CD	25N20		62.24	19CD		184CD1	54.10
19CD	25N20		74.29	19CD	25N20	19CG	14.62
184CE2	25N20	19NE2	98.89	184CE2		184CZ2	20.04
184CE2	25N20	162CE1	56.95	134CE2	25N20	184CD1	30.00

		T.	ABLE XII		
184CE2	25N20 162ND	1 68.56	184CE2	25N20 19CG	76.21
19NE2	25N20 162CE	1 69.93	19NE2	25N20 184CD1	73.25
19NE2	25N20 162ND	77.40	19NE2	25N20 19CG	29.26
184CZ2	25N20 162CE	1 57.21	184CZ2	25N20 184CD1	49.98
184CZ2	25N20 162ND	1 63.26	184CZ2	25N20 19CG	95.28
162CE1	25N20 184CD	1 61.93	162CE1	25N20 162ND1	16.79
162CE1	25N20 19CG	72.21	184CD1	25N20 162ND1	78.20
184CD1	25N20 19CG	47.15	162ND1	25N20 19CG	86.13
25SG	25C21 25CB	34.64	25SG	25C21 25N	72.39
25SG	25C21 25CA	49.66	25 <b>S</b> G	25C21 162ND1	48.43
25SG	25C21 19NE	93.26	25SG	25C21 26N	53.17
25 <b>S</b> G	25C21 1610	82.24	25 <i>S</i> G	25C21 25C	43.41
25SG	25C21 24C	80.54	25 <i>S</i> G	25C21 162CE1	48.10
25SG	25C21 19OE	L 73.78	25 <i>S</i> G	25C21 24CA	98.55
25 <b>S</b> G	25C21 163N	27.76	25SG	25C21 162CA	50.66
25SG	25C21 19CD	82.46	25SG	25C21 26CD1	90.92
25CB	25C21 25N	45.83	25CB	25C21 25CA	23.90
25CB	25C21 162ND1	58.24	25CB	25C21 19NE2	60.12
25CB	25C21 26N	51.36	25CB	25C21 24N	83.08
25CB	25C21 25C	33.89	25CB	25C21 24C	55.62
25CB	25C21 162CE1	46.38	25CB	25C21 19OE1	47.87
25CB	25C21 24CA	73.29	25CB	25C21 163N	62.27
25CB	25C21 162CA	80.47	25CB	25C21 19CD	51.61
25CB	25C21 26CD1	90.22	25N	25C21 25CA	23.31
25N	25C21 23O	62.57	25N	25C21 23C	56.71
25 <b>N</b>	25C21 23CA	72.88	25N	25C21 19NE2	53.88
25N	25C21 26N	39.27	25 <b>N</b>	25C21 24N	40.31
25N	25C21 25C	33.06	25N	25C21 24C	9.89
25N	25C21 162CE1	88.84	25N	25C21 19OE1	66.88
25N	25C21 24CA	27.62	25N	25C21 163N	97.26
25N	25C21 19CD	57.91	25N	25C21 26CD1	57.03
25CA	25C21 230	85.08	25CA	25C21 23C	79.97
25CA	25C21 162ND1	82.14	25CA	25C21 23CA	94.92
25CA	25C21 19NE2	59.55	25CA	25C21 26N	34.96
25CA	25C21 24N	63.21	25CA	25C21 25C	19.25
25CA	25C21 24C	32.60	25CA	25C21 162CE1	69.83
25CA	25C21 19OE1	60.15	25CA	25C21 24CA	50.80
25CA	25C21 163N	75.98	25CA	25C21 19CD	57.10
25CA	25C21 26CD1	68.81	230	25C21 23C	18.23
230	25C21 23CA	34.62	230	25C21 19NE2	81.51
230	25C21 26N	74.79	230	25C21 24N	31.37
230	25C21 25C	84.83	230	25C21 24C	52.77
230	25C21 24CA	35.38		25C21 19CD	94.62
230	25C21 26CD1	42.49	23C	25C21 23CA	21.54

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23C	25C21	19NE2	63.77	BLE XII	25C21	26N	80.57
23C	25C21	24N	17.67	23C	25C21	25C	85.73
23C	25C21	24C	48.08	23C	25C21	190E1	91.44
23C	25C21	24CA	30.02	23C	25C21	19CD	77.14
23C	25C21	26CD1	56.82	162ND1	25C21	19NE2	82.13
162ND1	25C21	1610	66.39	162ND1	25C21	25C	86.84
162ND1	25C21	162CE1	16.39	162ND1	25C21	190E1	54.45
162ND1	25C21		49.87	162ND1	25C21	162CA	38.64
162ND1	25C21	19CD	68.76	23CA	25C21	19NE2	59.28
23CA	25C21	24N	33.06	23CA	25C21	24C	65.84
23CA	25C21	190E1	85.51	23CA	25C21	24CA	49.18
23CA	25C21	19CD	72.54	23CA	25C21	26CD1	76.75
19NE2	25C21	26N	91.47	19NE2	25C21	24N	51.72
19NE2	25C21	25C	78.62	19NE2	25C21	24C	57.38
19NE2	25C21	162CE1	66.80	19NE2	25C21	190E1	27.71
19NE2	25C21	24CA	59.21	19NE2	25C21	19CD	13.58
26N	25C21	24N	70.38	26N	25C21	25C	17.60
26N	25C21	24C	40.64	26N	25C21	162CE1	94.63
26N	25C21	190E1	94.99	26N	25C21	24CA	53.42
26N	25C21	163N	67.59	26N	25C21	162CA	96.43
26N	25C21	19CD	91.55	26N	25C21	26CD1	40.50
24N	25C21	25C	71.62	24N	25C21	24C	32.79
24N	25C21	190E1	78.31	24N	25C21	24CA	16.97
24N	25C21	19CD	64.11	24N	25C21	26CD1	58.21
1610	25C21	162CE1	82.76	1610	25C21	163N	55.62
1610	25C21	162CA	33.90	25C	25C21	24C	38.87
25C	25C21	162CE1	78.56	25C	25C21	190E1	78.01
25C	25C21	24CA	55.90	25C	25C21	163N	64.97
25C	25C21	162CA	92.64	25C	25C21	19CD	76.23
25C	25C21	26CD1	56.82	24C	25C21	162CE1	98.56
24C	25C21	190E1	74.12	24C	25C21	24CA	18.29
24C	25C21	19CD	63.65	24C	25C21	26CD1	49.92
162CE1	25C21	190E1	39.16	162CE1	25C21	163N	58.96
162CE1	25C21	162CA	54.05	162CE1	25C21	19CD	53.25
190E1	25C21	24CA	82.23	190E1	25C21	163N	94.60
190E1	25C21	162CA	93.01	190E1	25C21	19CD	14.34
24CA	25C21	19CD	69.26	24CA	25C21	26CD1	46.30
163N	25C21	162CA	28.85	163N	25C21	26CD1	94.44
25SG	25022	25N	77.15	25 <b>\$</b> G	25022	25CB	38.56
25SG	25022	25CA	56.15	25 <b>S</b> G	25022	24C	89.30
25SG	25022	190E1	86.57	25 <i>S</i> G	25022		43.24
25 <b>S</b> G	25022	25C	48.29	25 <b>S</b> G	25022	26N	51.24
25SG	25022	162CE1	50.40	259G	25022	240	83.26
25N	25022	25CB	51.87	25N	25022	19NE2	73.58

			T	ABLE XII			
25N	25022		71.81	25N	25022	23CA	95.33
25N	25022		24.89	25N	25022	24N	52.99
25N	25022		73.36	25N	25022	24C	13.57
25N	25022	19CD	75.82	25N	25022	190E1	83.34
25N	25022	24CA	35.40	25N	25022	220	81.27
25N	25022	25C	29.02	25N	25022	26N	34.61
25N	25022	23N	91.35	25 <b>N</b>	25022	162CE1	94.53
25N	25022	22C	85.32	25 <b>N</b>	25022	240	6.50
25CB	25022	19NE2	77.77	25CB	25022	25CA	27.03
25CB	25022	24C	65.38	25CB	25022	19CD	66.62
25CB	25022	190E1	58.82	25CB	25022	24CA	87.22
25CB	25022	162ND1	52.46	25CB	25022	25C	32.94
25CB	25022	26N	48.44	25CB	25022	162CE1	44.82
25CB	25022	240	58.22	19NE2	25022	23C	87.29
19NE2	25022	23CA	80.68	19NE2	25022	25CA	75.11
19NE2	25022	24N	70.11	19NE2	25022	24C	73.32
19NE2	25022	19CD	15.53	19NE2	25022	190E1	32.76
19NE2	25022	24CA	76.24	19NE2	25022	220	41.66
19NE2	25022	162ND1	90.29	19NE2	25022	25C	91.32
19NE2	25022	23N	70.60	19NE2	25022	162CE1	74.46
19NE2	25022	22C	55.07	19NE2	25022	240	72.58
23C 23C	25022	23CA	27.25	23C	25022	25CA	96.67
23C	25022	24N	23.18	23C	25022	230	20.52
23C	25022	24C	58.32	23C	25022	24CA	36.50
23C	25022	220	49.90	23C	25022	25C	95.16
23C	25022 25022	26N	86.14	23C	25022	23N	30.10
23CA	25022 25022	22C	39.20	23C	25022	240	65.47
23CA	25022 25022	24N 24C	42.37	23CA	25022	230	41.32
23CA	25022	24CA	81.90	23CA	25022	19CD	95.72
23CA	25022	24CA 23N	60.58	23 <b>CA</b>	25022	220	39.10
23CA	25022	240	10.12	23CA	25022	22C	25.72
25CA ·	25022	230	88.84	25CA	25022	24N	77.38
25CA	25022	19CD	96.24 70.44	25CA	25022	24C	38.44
25CA	25022	24CA	60.28	25CA	25022	190E1	70.91
25CA		162ND1	79.49	25CA	25022	220	98.08
25CA	25022	26N	32.82	25CA	25022	25C	15.40
25CA	25022	240	31.31	25CA	25022		70.84
24N	25022	24C	39.66	24N	25022	230	38.32
24N	25022	24CA	19.43	24N	25022	19CD	83.95
24N	25022	25C	80.28	24N	25022	220	42.54
24N	25022	230 23N	39.05	24N	25022	26N	76.76
24N	25022	240	46.50	24N	25022	22C	38.22
230	25022	24CA	42.50	230	25022	24C	61.42
		2 TCA	-2.50	230	25022	220	70.35

			T	ABLE XII			
230	25022	25C	89.16	230	25022	26N	75.88
230	25022	23N	47.63	230	25022	22C	59.10
230	25022	240	68.00	24C	25022	19CD	79.18
24C	25022	190E1	89.99	24C	25022	24CA	21.85
24C	25022	220	71.83	24C	25022	25C	41.07
24C	25022	26N	42.17	24C	25022	23N	78.42
24C	25022	22C	73.88	24C	25022	240	7.15
19CD	25022	190E1	17.46	19CD	25022	24CA	86.98
19CD	25022	220	56.95	19CD	25022	162ND1	74.94
19CD	25022	25C	86.74	19CD	25022	23N	85.73
19CD	25022 1	62CE1	58.96	19CD	25022	22C	70.31
19CD	25022	240	76.60	190E1	25022	220	73.26
190E1	25022 1	62ND1	57.54	190E1	25022	25C	85.88
190E1		62CE1	41.81	190E1	25022	22C	86.38
190E1		240	85.82	24CA	25022	220	59.15
24CA		25C	61.20	24CA	25022	26N	57.49
24CA		23N	58.31	24CA	25022	22C	57.09
24CA		240	29.00	220	25022	23N	28.99
220		22C	13.41	220	25022	240	76.18
162ND1		25C	81.58	162ND1	25022	26N	91.32
162ND1		62CE1	16.64	25C	25022	26N	17.25
25C		62CE1	77.24	25C	25022	240	34.98
26N		62CE1	90.62	2 <b>6N</b>	25022	240	38.49
23N		22C	15.60	23N	25022	240	84.97
22C		240	79.45	1600	25C23	160CB	36.76
670H		67CE1	30.78	1600	25C24	160CB	45.57
1600		60C	10.69	1600	25C24	160CA	28.84
1600		50N	35.35	160CB	25C24	160C	34.90
160CB		09CD2	74.30	160CB	25C24	160CA	18.76
160CB		50N	30.63	160C	25C24	209CD2	96.30
160C		SOCA	18.91	160C	25C24	160N	29.27
67CE1		09CD2	57.14	67CE1	25C24	670H	30.07
209CD2		50CA	90.67	209CD2	25C24	670H	85.98
160CA	25C24 16		17.56	1600	25C25		5.82
1600	25C25 16		36.39	160C	25C25	160CB	31.49
67CE1		57OH	30.77	67CE1	25C25	67CZ	16.04
670H		7CZ	16.48	670H	25C26		31.37
670H		7CZ	1.6.61	67CE1	25C26		17.15
670H		7CZ	14.42	670H		67CE1	29.64
67CZ		7CE1	16.82	67CE1	25C27		94.55
670H		7CE1	29.88	670H	25C28	67CZ	13.63
67CE1		7CZ	16.55	2750H2	25030		92.49
1600		1CA	34.81	1600	25C31		41.59
1600	25C31 16	10	73.41	1600	25 <b>C</b> 31	161C	57.25

			Т	ABLE XII			
1600	25C31	160C	11.02	1600	25C3	l 161N	26.54
1600	25C31	161CB	50.31	161CA	25C3		34.32
161CA	25C31	. 161C	20.25	161CA	25C31		31.39
161CA	25C31	. 161N	15.77	161CA	25C31		16.64
1610	25C31	2750H2	83.79	1610	25C31		16.21
1610	25C31	160C	62.41	1610	25C31		47.12
1610	25C31	161CB	38.54	2750H2	25C31		97.89
2750H2	25C31	161CB	91.47	161C	25C31		46.24
161C	25C31	161N	31.16	161C	25C31		30.39
160C	25C31	161N	15.86	160C	25C31		42.23
161N	25C31		29.11	1600	25032		58.97
1600	25032	161C	79.84	1600	25032		17.38
1600	25032	161N	38.10	1600	25032		67.22
1600	25032	162N	74.42	1600	25032	160CA	13.85
161CA	25032	1610	45.01	161CA	25032	161C	27.02
161CA	25032	160C	43.17	161CA	25032	161N	22.39
161CA	25032	161CB	19.73	161CA	25032	162N	31.22
161CA	25032	160CA	49.48	1610	25032	161C	21.40
1610	25032	160C	83.84	1610	25032	161N	63.21
1610	25032	161CB	49.30	1610	25032	162N	29.81
1610	25032	160CA	88.75	1610	25032	2750H2	81.22
161C	25032	160C	62.49	161C	25032	161N	42.18
161C		161CB	39.00	161C	25032	162N	12.22
161C	25032	160CA	67.34	160C	25032	161N	21.25
160C	25032	161CB	55.00	160C	25032	162N	57.22
160C	25032	160CA	6.68	161N	25032	161CB	37.70
161N	25032	162N	39.25	161N	25032	160CA	27.25
161CB	25032	162N	47.45	161CB	25032	160CA	61.67
161CB		275OH2	93.88	162N	25032	160CA	61.00
1610	25C33	161C	15.17	1610	25C33	2750H2	81.48
1610	25C33	161CA	<b>29.1</b> 9	660	25C33	66N	37.60
660	25C33	65CA	66.72	161C	25C33	2750H2	91.71
161C	25C33	161CA	18.22	6 <b>6N</b>	25C33	2750H2	84.36
66N	25C33	65CA	<b>29.</b> 78	2750H2	25C33	161CA	86.95
275OH2	25C33	65CA	56.91	660	25C34	66C	9.95
660	25C34	66N	42.43	660	25C34	26CB	47.74
660	25C34	66CA	25.52	660	25C34	67N	13.89
66Q	25C34	26CG	42.21	660	25C34	163CB	91.47
66C	25C34	66N	36.15	66C	25C34	26CB	56.11
66C	25C34	66CA	18.24	65C	25C34	67 <b>N</b>	11.90
66C	25C34	26CG	48.09	6 <b>6N</b>	25C34	26CB	65.07
66N	25C34	66CA	17.98	66N	25C34	67N	47.31
66N	25C34	26CG	48.82	26CB	25C34	66CA	60.10
26CB	25C34	67N	60.76	26CB	25C34	26CG	17.03

				ABLE XII			
26CB	25C34		52.12	66CA	25C34	67N	29.36
66CA	25C34		46.86	1610	25C34	163CB	78.29
67N	25C34		56.10	67N	25C34	163CB	99.58
26CG	25C34	163CB	68.12	660	25C35	163CB	89.09
660	25C35	66C	7.53	660	25C35	67CD1	63.13
660	25C35	68SD	68.36	660	25C35	26CB	39.72
209CD2	25C35	134CB	51.38	209CD2	25C35	67CD1	51.16
209CD2	25C35	68SD	69.83	209CD2	25C35	1600	84.17
134CB	25C35	163CB	64.65	134CB	25C35	163N	61.10
134CB	25C35		66.71	134CB	25C35	1600	74.73
134CB	25C35	1610	92.07	163CB	25C35	66C	96.13
163CB	25C35	163N	30.24	163CB	25C35	68SD	45.97
163CB	25C35	26CB	50.88	163CB	25C35	1610	80.11
66C	25C35		56.50	66C	25C35	68SD	71.94
66C	25C35		47.21	67CD1	25C35	68SD	80.40
67CD1	25C35		97.36	163N	25C35	68SD	74.50
163N	25C35		89.86	163N	25C35	26CB	73.24
163N	25C35		52.41	68SD	25C35	26CB	52.35
1600	25C35		56.45	26CB	25C35	1610	98.49
161C	25C36		18.57	161C	25C36	162N	20.01
161C	25C36		62.57	161C	25C36	163N	67.45
161C	25C36		21.83	161C	25C36	162C	54.00
161C	25C36		53.82	161C	25C36	161N	36.26
161C	25C36		34.07	161C	25C36	163CA	85.21
161C	25C36		99.05	161C		1620	59.74
161C	25C36		80.93	161C		134CA	96.31
1610	25C36		33.64	1610	25C36	1600	74.48
1610	25C36		66.90	1610	25C36	161CA	34.96
1610	25C36	162C	59.06	1610	25C36	160C	69.44
1610	25C36	161N	53.14	1610	25C36	162CA	38.64
1610	25C36	163CA	85.32	1610	25C36	163CB	94.16
1610	25C36	1620	69.06	1610	25C36	160CB	98.55
162N		134CB	89.78	162N		1600	71.09
162N	25C36		52.42	162N	25C36		35.49
162N	25C36		36.12	162N	25C36		57.50
162N	25C36		39.78	162N	25C36		19.47
162N		163CA	68.57	162N	25C36		85.10
162N	25C36		39.89	162N		160CB	76.82
162N	25C36		77.09	134CB		1600	94.70
134CB	25C36		73.72	134CB	25C36		70.42
134CB	25C36		84.32	134CB		161N	89.08
134CB		162CA	87.63	134CB		163CA	61.39
134CB	25C36	163CB	69.65	134CB		1620	57.00
134CB	25C36	209CD2	50.87	134CB	25C36	160CB	57.37

		1	ABLE XII		
134CB	25C36 134CA	14.58	1600	25C36 161CA	40.90
1600	25C36 160C	17.14	1600	25C36 161N	31.70
1600	25C36 162CA	90.56	1600	25C36 162O	96.89
1600	25C36 209CD2	91.01	1600	25C36 160CB	37.47
1600	25C36 134CA	97.11	163N	25C36 161CA	87.43
163N	25C36 162C	18.84	163N	25C36 161N	90.13
163N	25C36 162CA	33.77	163N	25C36 163CA	18.45
163N	25C36 163CB	32.69	163N	25C36 1620	29.16
163N	25C36 134CA	60.25	161CA	25C36 162C	71.59
161CA	25C36 160C	34.60	161CA	25C36 161N	20.18
161CA	25C36 162CA	53.69	161CA	25C36 1620	73.10
161CA	25C36 160CB	64.88	161CA	25C36 134CA	99.89
162C	25C36 160C	86.67	162C	25C36 161N	71.72
162C	25C36 162CA	20.71	162C	25C36 163CA	32.55
162C	25C36 163CB	50.28	162C	25C36 1620	14.79
162C	25C36 160CB	91.85	162C	25C36 134CA	55.85
160C	25C36 161N	18.17	160C	25C36 162CA	76.54
160C	25C36 1620	79.80	160C	25C36 209CD2	94.97
160C	25C36 160CB	31.31	160C	25C36 134CA	83.64
161N	25C36 162CA	59.12	161N	25C36 1620	67.84
161N	25C36 160CB	45.47	161N	25C36 134CA	83.93
162CA	25C36 163CA	51.16	162CA	25C36 163CB	66.24
162CA	25C36 162O	30.96	162CA	25C36 160CB	91.92
162CA	25C36 134CA	73.28	163CA	25C36 163CB	19.41
163CA	25C36 1620	35.58	163CA	25C36 134CA	50.10
163CA	25C36 660	93.58	163CB	25C36 1620	54.97
163CB	25C36 209CD2	96.16	163CB	25C36 134CA	62.02
163CB	25C36 660	74.19	1620	25C36 160CB	79.17
1620	25C36 134CA	42.49	209CD2	25C36 160CB	65.91
209CD2	25C36 134CA	65.34	209CD2	25C36 660	80.01
160CB	25C36 134CA	61.85	209CD2	25C37 67CD1	67.58
209CD2	25C37 67CE1	72.61	209CD2	25C37 67CG	78.13
209CD2	25C37 1600	97.02	209CD2	25C37 67CZ	85.11
209CD2	25C37 134CB	51.48	209CD2	25C37 67CA	84.67
209CD2	25C37 209CG	2.50	67CD1	25C37 67CE1	21.97
67CD1	25C37 660	73.49	67CD1	25C37 67CG	14.24
67CD1	25C37 67CZ	29.56	67CD1	25C37 66C	63.11
67CD1	25C37 67CA	43.36	67CD1	25C37 209CG	66.09
67CE1	25C37 660	88.74	67CE1	25C37 67CG	32.18
67CE1	25C37 67CZ	12.63	67CE1	25C37 66C	77.14
67CE1	25C37 67CA	63.67	67CE1	25C37 209CG	72.01
660	25C37 67CG	59.39	660	25C37 67CZ	84.53
660	25C37 66C	11.85	660	25C37 67CA	35.08
67CG	25C37 67CZ	34.96	67CG	25C37 66C	48.87

			т	ABLE XII			
67CG	25C37	7 67CA	31.56	67CG	25C3	7 209CG	76.30
1600	25C37	134CB	73.49	1600	25C3		99.48
67CZ	25C37	66C	72.68	67CZ	25C31		65.61
67CZ	25C37	209CG	84.57	66C	25C3		29.50
134CB	25C37	209CG	52.27	67CA	25C31		82.22
65CA	25C38	66N	35.46	65CA	25C38		74.46
65CA	25C38	26CD1	58.84	65CA	25C38		67.35
65CA	25C38	65C	19.07	65CA	25C38		55.20
65CA	25C38	26CB	86.32	65CA	25C38		69.11
65CA	25C38	640	33.75	65CA	25C38		11.94
1610	25C38	25 <i>S</i> G	65.22	1610	25C38		84.21
1610	25C38	161C	10.24	6 <b>6N</b>	25C38		39.10
66N	25C38	26CD1	49.39	6 <b>6N</b>	25C38	2750H2	95.80
66N	25C38	65C	16.72	66N	25C38	230	75.10
66N	25C38	26CB	62.01	66N	25C38	26CG	49.74
66N	25C38	640	67.28	6 <b>6N</b>	25C38	65N	46.94
660	25C38	26CD1	55.62	660	25C38	65C	55.45
660	25C38	230	98.46	660	25C38	26CB	41.40
660	25C38	26CG	43.32	660	25C38	65N	85.60
26CD1	25C38	25SG	77.15	26CD1	25C38	65C	49.46
26CD1	25C38	230	44.41	26CD1	25C38	26CB	32.56
26CD1	25C38	26CG	15.85	26CD1	25C38	640	87.10
26CD1	25C38	65N	61.51	25SG	25C38	230	70.52
25 <b>S</b> G	25C38	26CB	68.80	25SG	25C38	26CG	76.92
25SG	25C38	161C	68.97	2750н2	25C38	65C	83.85
2750H2	25C38	230	87.47	2750H2	25C38	640	35.10
2750H2	25C38	161C	89.52	2750H2	25C38	65N	61.32
65C	25C38	230	62.91	65C	25C38	26CB	71.13
65C	25C38	26CG	55.52	65C	25C38	640	52.20
65C	25C38	65N	30.26	230	25C38	26CB	73.10
230	25C38	26CG	59.68	230	25C38	640	61.47
230	25C38	65N	47.46	26CB	25C38	26CG	17.91
26CB	25C38	65N	91.96	26CG	25C38	65N	74.13
640 65CA	25C38	65N	26.26	65CA	25039	66N	45.40
65CA	25039	65C	23.65	65CA	25039	2750H2	89.27
65CA	25039	640	46.55	65CA	25039	65N	17.11
65CA	25039	26CD1	69.73	65CA	25039	230	68:59
65CA	25039	64C	33.21	65CA	25039	660	84.46
65CA	25039	66CA	54.04	65CA	25039	650	20.83
65CA	25039	26CG	77.43	65CA	25039	26NE1	55.20
66N	25039	66C	71.68	66N	25039	65C	22.01
	25039	640	89.22	66N	25039	65N	62.07
66N	25039	26CD1	54.96	66N	25039	230	90.19
66N	25039	64C	78.37	66N	25039	66C	39.27

c c			7	ABLE XII			
66N	25039		8.76	6 <b>6N</b>	2503	9 650	25.54
66N	25039	26CG	51.40	66N	2503	9 26NE1	49.00
66N	25039	66C	26.33	65C	2503		69.28
65C	25039	65N	40.08	65C	25039		57.43
65C	25039	230	76.91	65C	25039		56.85
65C	25039	660	60.83	65C	25039	_	30.50
65C	25039	650	3.97	65C	25039		60.36
65C	25039	26NE1	45.73	65C	25039		48.13
2750H2	25039	640	44.70	2750н2			79.36
2750H2	25039	64C	61.39	2750н2		1610	78.76
640	25039	65N	34.67	640	25039		76.69
640	25039	64C	16.94	640	25039		97.89
640	25039	650	67.08	640	25039		93.96
65 <b>N</b>	25039	26CD1	75.00	65N	25039		58.94
65N	25039	64C	18.53	65N	25039		
65N	25039	650	36.76	65N	25039		70.56
65N	25039	26NE1	60.15	65N	25039	66C	85.84
26CD1	25039	230	49.08	26CD1	25039	64C	88.15
26CD1	25039	660	56.07	26CD1	25039	66CA	91.52
26CD1	25039	650	56.44	26CD1	25039	26CG	53.07
26CD1	25039	26NE1	14.91	26CD1	25039	66C	14.63 54.99
230	25039	64C	64.20	230	25039	66CA	93.33
230	25039	650	73.30	230	25039	26CG	63.07
230	25039	26NE1	44.74	64C	25039	66CA	
64C	25039	650	53.98	64C	25039	26NE1	87.09 77.02
660	25039	66CA	30.52	660	25039	650	63.93
660	25039	26CG	42.81	660	25039	26NE1	62.05
660	25039	66C	13.06	66CA	25039	650	33.85
66CA	25039	26CG	47.00	66CA	25039	26NE1	49.86
66CA	25039	66C	17.64	650	25039	26CG	60.51
650	25039	26NE1	44.00	650	25039	66C	51.39
26CG	25039	26NE1	26.93	26CG	25039	66C	44.17
26NE1	25039	66C	57.23	25 <i>S</i> G	25 <b>N4</b> 0	1610	83.77
25 <b>S</b> G	25N40	26CD1	96.23	25 <i>S</i> G	25N40	26N	54.32
25 <b>S</b> G	25N40	230	88.72	25 <b>S</b> G	25N40	26CB	85.40
25SG	25N40 1	.63N	48.48	25 <i>S</i> G	25N40	25CB	12.78
25SG	25N40	26CG	93.55	25 <b>S</b> G		161C	83.44
25 <b>S</b> G		25N	43.46	25 <b>S</b> G	25N40		55.83
25SG		26CA	67.26	25 <b>S</b> G	25N40	23C	
25SG		63CB	56.68	25 <i>S</i> G	25N40	25CA	77.46
25 <b>S</b> G		25C	40.47	1610		23CA 163N	30.38
1610		25CB	96.42	1510		161C	62.30
1610		62CA	34.15	1610		163CB	9.97
26CD1		26N	45.97	26CD1	25N40 25N40		90.03
<del>-</del>			-3.31	2 OCDI	23M40	230	47.29

2601	21 25242	7	TABLE XI	Ī	
26CI		34.59	26CD		83.86
26CI	1	16.42	26CD		56.97
26CI		53.52	26CD	1 25 <b>N4</b> 0 660	51.90
26CD		42.89	26CD		44.25
26CD		56.28	26CD1		84.60
26CD		66.03	26CD1		58.27
26N	25N40 230	70.59	26N	25N40 26CB	33.25
26N	25N40 163N	70.68	26N	25N40 25CB	44.21
26N	25N40 26CG	39.40	26N	25N40 25N	34.54
26N	25N40 65CA	99.36	26N	25N40 660	73.26
26N	25N40 162CA	97.09	26N	25N40 26CA	16.76
26N	25N40 66N	85.93	26N	25N40 23C	69.38
26N	25N40 163CB	48.13	26N	25N40 25CA	29.29
26N	25N40 25C	13.85	230	25N40 26CB	79.60
230	25N40 25CB	77.95	230	25N40 26CG	63.18
230	25N40 25N	49.38	230	25N40 65CA	52.49
230	25N40 660	94.81	230	25N40 26CA	79.46
230	25N40 66N	70.07	230	25N40 23C	13.03
230	25N40 25CA	66.29	230	25N40 25C	73.80
26CB	25N40 163N	81.30	26CB	25N40 25CB	76.86
26CB	25N40 26CG	18.52	26CB	25N40 25N	64.46
26CB	25N40 65CA	80.54	26CB	25N40 660	40.01
26CB	25N40 26CA	18.15	26CB	25N40 66N	57.97
26CB	25N40 23C	85.39	26CB	25N40 163CB	51.83
26CB	25N40 25CA	62.52	26CB	25N40 25C	46.36
163N	25N40 25CB	57.29	163N	25N40 26CG	99.16
163N	25N40 161C	55.09	163N	25N40 25N	84.86
163N	25N40 660	96.57	163N	25N40 162CA	29.80
163N	25N40 26CA	70.11	163N	25N40 163CB	29.52
163N	25N40 25CA		163N	25N40 25C	62.99
25CB	25N40 26CG	82.58	25CB	25N40 161C	96.21
25CB	25N40 25N	30.92	25CB	25N40 162CA	68.31
25CB	25N40 26CA	58.91	25CB	25N40 23C	67.68
25CB 25CB	25N40 163CB	58.44	25CB	25N40 25CA	17.83
25CB 26CG	25N40 25C	30.57	26CG	25N40 25N	61.46
	25N40 65CA	64.22	26CG	25N40 660	41.69
26CG	25N40 26CA	30.63	26CG	25N40 66N	46.53
26CG	25N40 23C	71.01	26CG	25N40 163CB	69.91
26CG	25N40 25CA	65.57	26CG	25N40 25C	53.15
161C	25N40 162CA			25N40 163CB	81.51
25N	25N40 65CA	97.53		25N40 162CA	99.23
25N	25N40 26CA	50.89		25N40 23C	41.97
25N	25N40 163CB	74.65		25N40 25CA	16.92
25N	25N40 25C	28.88	65CA	25N40 660	62.01

			T	ABLE XII			
65CA	25 <b>N4</b> (		94.78	65CA	25 <b>N4</b> 0	66N	28.94
65CA	25N40		64.73	660	25N40	26CA	57.46
660	25 <b>N4</b> 0	66N	33.09	660	25N40	163CB	72.28
660	25 <b>N4</b> 0	25C	86.11	162CA	25 <b>N4</b> 0	26CA	99.40
162CA	25 <b>N4</b> 0	163CB	59.22	162CA	25N40	25CA	84.50
162CA	25 <b>N4</b> 0		86.43	26CA	25 <b>N4</b> 0	66N	75.24
26CA	25 <b>N4</b> 0		81.32	26CA	25N40	163CB	42.44
26CA	25 <b>N4</b> 0		45.63	26CA	25N40	25C	28.66
66N	25 <b>N4</b> 0	23C	83.06	66N	25 <b>N4</b> 0	25C	99.63
23C	25 <b>N4</b> 0		58.57	23C	25 <b>N4</b> 0	25C	69.36
163CB	25 <b>N4</b> 0		59.45	163CB	25 <b>N4</b> 0	25C	46.39
25CA	25 <b>N4</b> 0	25C	17.68	25SG	25C41	25N	62.70
25 <b>S</b> G	25C41	26N	62.65	25SG	25C41	25CB	23.96
25SG	25C41	25CA	43.98	25SG	25C41	24N	98.96
25\$G	25C41	24C	76.68	25 <i>S</i> G	25C41	25C	48.95
25 <i>S</i> G	25C41	26CB	86.53	25SG	25C41	24CA	95.08
25 <b>S</b> G	25C41	1610	70.41	25SG	25C41	26CA	70.68
230	25C41	25N	69.35	230	25C41	23C	19.72
230	25C41	26CD1	58.41	230	25C41	26N	90.56
230	25C41	25 <b>CA</b>	89.48	230	25C41	23CA	34.94
230	25C41	24N	31.63	230	25C41	24C	57.25
230	25C41	25C	94.72	230	25C41	26CB	91.35
230	25C41	26CG	73.21	230	25C41	24CA	37.60
230	25C41	65CA	57.07	230	25C41	26NE1	46.32
230	25C41	26CA	94.04	25N	25C41	23C	58.58
25N	25C41	26CD1	71.53	25N	25C41	26N	42.84
25N	25C41	25CB	39.04	25 <b>N</b>	25C41	25CA	20.25
25N	25C41	23CA	71.04	25N	25C41	24N	41.24
25N	25C41	24C	14.76	25 <b>N</b>	25C41	25C	33.26
25N	25C41	26CB	72.91	25 <b>N</b>	25C41	26CG	71.59
25N	25C41	24CA	32.43	2 <b>5N</b>	25C41	26NE1	71.77
25N	25C41	26CA	57.10	23C	25C41	26CD1	72.85
23C	25C41	26N	90.94	23C	25C41	25CB	91.53
23C	25C41	25CA	78.46	23C	25C41	23CA	20.98
23C	25C41	24N	17.34	2 <b>3</b> C	25C41	24C	50.23
23C	25C41	25C	89.47	23C	25C41	26CG	86.15
23C	25041	24CA	32.48	230	25C41	65CA	75.78
23C	25C41	26NE1	61.91	23C	25C41	26CA	99.05
26CD1	25C41	26N	51.41	26CD1	25C41	25CA	79.66
26CD1	25C41	23CA	92.59	26CD1	25041	24N	70.00
26CD1	25C41	24C	58.88	26CD1	25C41	25C	66.43
26CD1	25C41	26CB	34.11	26CD1	25C41	26CG	15.31
26CD1	25C41	24CA	54.83	26CD1		65CA	55.19
26CD1	25C41	26NE1	12.42	26CD1	25C41	26CA	44.47

		1	ABLE XII			
26N	25C41 25CB	51.65	26N	25C41	25CA	35.77
26N	25C41 24N	75.80	26N	25C41		42.85
26N	25C41 25C	16.65	26N	25C41		33.09
26N	25C41 26CG	41.38	26N	25C41		58.47
26N	25C41 26NE1	60.33	26N	25C41		14.86
25CB	25C41 25CA	22.08	25CB	25C41		94.98
25CB	25C41 24N	75.58	25CB	25C41		53.42
25CB	25C41 25C	35.13	25CB	25C41		82.86
25CB	25C41 26CG	92.88	25CB	25C41	24CA	71.25
25CB	25C41 1610	93.68	25CB	25C41	26CA	64.25
25CA	25C41 23CA	88.69	25CA	25C41	24N	61.20
25CA	25C41 24C	32.93	25CA	25C41	25C	20.29
25CA	25C41 26CB	68.87	25CA	25C41	26CG	74.42
25CA	25C41 24CA	52.13	25CA	25C41	26NE1	83.73
25CA	25C41 26CA	50.39	23CA	25C41	24N	33.53
23CA	25C41 24C	66.67	23CA	25C41	24CA	51.61
23CA	25C41 65CA	80.98	23CA	25C41	26NE1	80.92
24N	25C41 24C	33.63	24N	25C41	25C	72.54
24N	25C41 26CB	93.33	24N	25C41	26CG	80.45
24N	25C41 24CA	18.67	24N	25C41	65CA	88.30
24N	25C41 26NE1	61.55	24N	25C41	26CA	85.94
24C	25C41 25C	39.29	24C	25C41	26CB	67.28
24C	25C41 26CG	61.65	24C	25C41	24CA	19.66
24C	25C41 26NE1	57.67	24C	25C41	26CA	54.90
25C	25C41 26CB	49.26	25C	25C41	26CG	57.75
25C	25C41 24CA	58.24	25C	25C41	26NE1	73.78
25C	25C41 26CA	30.51	26CB	25C41	26CG	18.80
26CB	25C41 24CA	74.83	26CB	25C41	65CA	78.82
26CB	25C41 26NE1	46.51	26CB	25C41	26CA	18.80
26CG	25C41 24CA	63.20	25CG	25C41	65CA	65.01
26CG	25C41 26NE1	27.72	26C <b>G</b>	25C41	26CA	31.31
24CA	25C41 65CA	87.25	24CA	25C41	26NE1	48.94
24CA	25C41 26CA	67.55	65CA	25C41	26NE1	49.22
65CA	25C41 26CA	95.86	26NE1	25C41	26CA	55.54
2750H2	25N42 1610	84.19	2750 <b>H2</b>	25N42	66N	91.61
2750H2	25N42 65CA	62.05	2750H2	25N42 1		94.32
2750H2	25N42 161CA		1.610	25N42 1	161C	13.86
1610	25N42 1600	61.38	1610	25N42 1	l61CA	29.35
66N	25N42 660	34.84	6 <b>6N</b>	25N42	65CA	29.86
660	25N42 65CA	63.11	161C	25N42 1	L600	47.97
161C	25N42 161CA	17.98	1600	25N42 1	L61CA	33.22

## TABLE XIII

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide.

Atom :	l Atom	2 Atom 3	Angle	Atom 1	Atom	2 Atom 3	Angle
184CB	25C1	1840	38.19	184CB	25C1	184CG	19.52
184CB	25C1	184CD2	33.79	184CB	25C1		43.09
184CB	25C1	188CD1	58.81	1840	25C1		56.83
1840	25C1	184CD2	71.93	1840	25C1	184CE3	78.59
1840	25C1	188CD1	69.71	184CG	25C1	184CD2	17.68
184CG	25C1	184CE3	32.58	184CG	25C1	188CD1	65.18
184CD2	25C1	184CE3	17.04	184CD2	25C1	188CD1	58.90
184CE3	25C1	188CD1	45.42	1840	25C2	184CB	44.21
1840	25C2	180D1	53.44	1840	25C2	184C	13.95
1840	25C2	184CA	33.42	1840	25C2	184CG	63.00
184CB	25C2	180D1	67.91	184CB	25C2	184C	34.94
184CB	25C2	184CA	20.32	184CB	25C2	184CG	19.43
180D1	25C2	184C	45.96	180D1	25C2	184CA	48.12
180D1	25C2	184CG	73.56	184C	25C2	184CA	20.19
184C	25C2	184CG	52.12	184CA	25C2	184CG	33.37
180D1	25C3	184CB	80.07	180D1	25C3	1840	59.68
180D1	25C3	184CA	57.15	180D1	25C3	184CG	89.66
180D1	25C3	184C	51.33	180D1	25C3	184CD1	82.13
180D1	25C3	18CG	11.24	180D1	25C3	18ND2	27.16
180D1	25C3	200	73.60	184CB	25C3	1840	44.39
184CB	25C3	184CA	22.92	184CB	25C3	184CG	22.24
184CB	25C3	184C	36.34	184CB	25C3	184CD1	37.05
184CB	25C3	18CG	84.56	184CB	25C3	18ND2	96.50
184CB	25C3	184CD2	33.32	1840	25C3	184CA	36.38
1840	25C3	184CG	66.41	1840	25C3	184C	16.67
1840	25C3	184CD1	77.34	1840	25C3	18CG	56.51
1840	25C3	18ND2	60.61	1840	25C3	184CD2	76.89
184CA	25C3	184CG	37.43	184CA	25C3	184C	21.29
184CA	25C3	184CD1	42.36	184CA	25C3	18CG	62.14
184CA	25C3	18ND2	75.37	184CA	25C3	184CD2	52.39
184CG	25C3	184C	56.26	184CG	25C3	184CD1	18.80

	10400	2502	1000		ABLE XIII			
	184CG	25C3		97.29		25C3	184CD2	15.81
	184CG	25C3		99.33	184C	25C3	184CD1	63.60
	184C	25C3		51.80	184C	25C3	18ND2	61.00
	184C	25C3	<del></del>	69.50	184CD1	25C3	18CG	91.92
	184CD1			27.91	184CD1	25C3	200	80.68
	18CG	25C3	18ND2	16.43	18CG	25C3	200	80.57
	18ND2	25C3	200	85.65	184CG	25C4	184CD1	22.16
	184CG	25C4		23.25	184CG	25C4	184NE1	33.38
	184CG	25C4	184CD2	20.04	184CG	25C4	184CA	36.56
	184CG	25C4	180D1	84.23	184CG	25C4	184CE2	31.43
	184CG	25C4	1840	59.49	184CG	25C4	184CE3	31.14
	184CG	25C4	184C	50.73	184CD1	25C4	184CB	40.60
:	184CD1	25C4	184NE1	19.22	184CD1	25C4	184CD2	33.07
	184CD1	25C4	184CA	43.38	184CD1	25C4	180D1	80.63
:	184CD1	25C4	184CE2	30.49	184CD1	25C4	200	96.73
:	184CD1	25C4	1840	72.55	184CD1	25C4	184CE3	47.65
•	L84CD1	25 <b>C4</b>	184C	60.18	184CB	25C4	184NE1	55.95
	L84CB	25C4	184CD2	38.71	184CB	25C4	184CA	20.96
1	L84CB	25 <b>C4</b>	180D1	69.40	184CB	25C4	184CE2	54.07
	L84CB	25C4	1840	36.44	184CB	25C4	184CE3	42.67
1	L84CB	25C4	184C	29.74	184NE1	25C4	184CD2	31.83
1	84NE1	25 <b>C4</b>	184CA	62.30	184NE1	25C4	180D1	97.90
1	84NE1	25 <b>C</b> 4	184CE2	18.20	184NE1	25C4	200	93.72
1	.84NE1	25C4	1840	90.55	184NE1	25C4	184CE3	45.60
1	84NE1	25C4	184C	78.96	184CD2	25C4	184CA	55.86
1	.84CD2	25C4	184CE2	18.77	184CD2	25C4	1840	74.24
1	.84CD2	25 <b>C4</b>	184CE3	14.78	184CD2	25C4	184C	68.30
1	.84CA	25C4	180D1	49.23	184CA	25C4	184CE2	67.15
1	84CA	25C4	1840	30.22	184CA	25C4	184CE3	62.73
1	84CA	25C4	184C	16.85	180D1	25C4	200	73.28
	180D1	25C4	1840	46.75	1.80D1	25C4	184C	42.18
1	84CE2	25C4	1840	90.51	184CE2	25C4	184CE3	29.58
1	84CE2	25C4	184C	82.12	1840	25C4	184CE3	73.98
1	840	25C4	184C	14.33	184CE3	25C4	184C	71.72
1	84CG	25C5	184CD2	22.73	184CG	25C5	184CD1	20.69
1	84CG	25C5	184CE2	35.01	184CG	25C5	184NE1	33.46
1	84CG	25C5	184CB	21.39	184CG	25C5	184CE3	38.46
1	84CG	25C5	184CZ2	50.35	184CG	25C5	184CZ3	51.39

1840	G 250	75 19402	•-	TABLE X			
184C						5 184CD1	34.47
1840						5 184NE1	
184CI			39.1			5 184CE3	19.36
184CI						5 184CZ3	29.20
184CI			51.3			5 184CE2	33.24
184CI	_		,			5 184CB	37.53
184CD	_						49.12
184CE				<del>-</del>			36.61
184CE						· - <del>-</del>	55.77
184CE						184CZ2	16.25
184NE			34.98				63.76
184NE			54.14				51.39
184NE			32.74				54.44
184CB	25C5		56.07	•	25C5		47.44
184CB	25C5		70.15		25C5	184CZ3	62.99
184CE3			16.19 15.81			184CZ2	37.12
184CZ2			29.69			184CA	62.73
184CZ3			77.85			184CA	79.60
184CD2			20.21			184CE3	19.98
184CD2			17.00	184CD2	25C6	184CB	36.83
184CD2		184CD1	27.90	184CD2	25C6	184CZ3	30.72
184CE3		184CB	46.74	184CE3	25C6	184CG	37.11
184CE3	25C6	184CZ3	16.21	184CE3	25C6	184CE2	31.55
184CG	25C6	184CB	20.58	184CE3 184CG	25C6	184CD1	47.70
184CG	25C6	184CZ3	50.48	184CG	25C6	184CE2	29.67
184CB	25C6	184CE2	49.78	184CB	25C6	184CD1	15.71
184CB	25C6	184CD1	32.55	184CE2	25C6	184CZ3	62.65
184CE2	25C6	184CD1	27.45	184CE2	25C6	184CZ3	34.33
200	25C7		65.72	200	25C6	184CD1	57.71
200	25C7	20N	42.52	200	25C7	20C	13.67
200	25C7	19CD	74.17	200	25C7	180D1	88.27
200	25 <b>C</b> 7	19NE2	65.37	200	25C7	20CA	32.34
200	25C7	19C	42.11	184CD1	25C7	190E1	87.32
184CD1	25 <b>C</b> 7	184NE1	21.55		25C7	19CG	67.76
184CD1	25C7	180D1	84.02	184CD1 184CD1		184CG	19.86
184CD1	25C7	184CE2	29.40	184CD1	25C7	19CD	60.66
184CD1	25C7	19NE2	73.46	184CD1		184CB	34.91
		-: <del></del>		TOWCDI	25 <b>C</b> 7	184CD2	28.08

			7	ABLE XIII			
184CD	1 25C	7 184CA	40.17			1830	43.72
184CD	1 25C	7 190E1	47.61	184CD1	. 25C7	19C	99.25
19CG	25C	7 184NE1	69.79	19CG	25C7	184CG	86.16
19CG	25C	7 20C	70.46	19CG	25 <b>C</b> 7	20N	50.72
19CG	25C7	7 180D1	75.31	19CG	25C7	19CD	20.77
19CG	25C7	7 20CA	67.17	19CG	25C7	184CE2	86.39
19CG	25C7	184CB	91.37	19CG	25C7	19NE2	32.57
19CG	25C7	184CD2	94.84	19CG	25C7	184CA	78.19
19CG	25C7	1830	44.08	19CG	25C7	190E1	28.49
19CG	25C7	19C	38.77	184NE1	25C7	184CG	34.11
184NE1	25C7	19CD	55.36	184NE1	25C7	184CE2	16.60
184NE1	25C7	184CB	52.97	184NE1	25C7	19NE2	62.84
184NE1	25C7	184CD2	29.31	184NE1	25C7	184CA	61.52
184NE1	25C7	1830	60.91	184NE1	25C7	190E1	42.84
184CG	25 <b>C</b> 7	180D1	81.09	184CG	25C7	19CD	80.51
184CG	25C7	184CE2	31.09	184CG	25C7	184CB	19.37
184CG	25C7	19NE2	92.98	184CG	25C7	184CD2	17.53
184CG	25C7	184CA	33.64	184CG	25C7	1830	53.42
184CG	25C7	190E1	67.47	20C	25C7	20N	35.03
20C	25C7	180D1	76.84	20C	25C7	19CD	82.78
20C	25C7	20CA	20.30	20C	25C7	19NE2	76.56
20C	25C7	1830	97.75	20C	25C7	190E1	95.52
20C	25C7	19C	38.85	20N	25C7	180D1	48.08
20N	25C7	19CD	70.24	20 <b>N</b>	25 <b>C</b> 7	20CA	19.43
20N	25 <b>C</b> 7	19NE2	74.34	20N	25C7	184CA	87.18
20N	25C7	1830	63.69	20N	25C7	190E1	79.16
20N	25C7	19C	12.46	180D1	25C7	19CD	94.22
180D1	25 <b>C7</b>	20CA	56.55	180D1	25C7	184CB	64.46
180D1	25C7	184CD2	98.02	180 <b>D1</b>	25C7	184CA	47.47
180D1	25C7	1830	48.16	180D1	25C7	190E1	93.80
180D1	25C7	19C	55.42	19CD	25 <b>C</b> 7	20CA	84.64
19CD	25C7	184CE2	71.23	19CD	25C7	184CB	91.94
19CD	25C7	19NE2	17.10	1.9CD	25C7	184CD2	83.77
19CD	25 <b>C</b> 7	184CA	84.58	19CD	25C7	1830	54.34
19CD	25C7	190E1	13.19	19CD	25C7	19C	57.87
20CA	25C7	19NE2	84.21	20CA	25 <b>C</b> 7	1830	82.52
SOCY	25C7	190E1	95.40	20CA	25C7	19C	28.93
184CE2	25C7	184CB	50:04	184CE2	25 <b>C</b> 7	19NE2	76.55

			T	ABLE XIII			
184CE2	25C7	184CD2	18.00	184CE2	25C7	184CA	64.00
184CE2	25C7	1830	72.92	184CE2	25C7	190E1	59.12
184CB	25C7	184CD2	34.04	184CB	25C7	184CA	19.42
184CB	25C7	1830	50.23	184CB	25C7	190E1	79.91
19NE2	25C7	184CD2	92.14	19NE2	25 <b>C</b> 7	1830	71.31
19NE2	25C7	190E1	27.20	19NE2	25C7	19C	62.12
184CD2	25C7	184CA	50.81	184CD2	2 <b>5</b> C7	1830	69.19
184CD2	25C7	190E1	70.74	184CA	25C7	1830	34.36
184CA	25C7	190E1	74.90	184CA	25C7	19C	87.64
1830	25C7	190E1	48.64	1830	25 <b>C</b> 7	19C	59.18
190E1	25C7	19C	67.25	200	2508	20C	10.26
200	2508	19CG	58.52	200	2508	19NE2	68.66
200	2508	19CD	71.25	200	2508	20CA	23.15
200	2508	20N	31.47	200	2508	21N	16.79
200	2508	21CA	28.59	20C	2508	19CG	64.70
20C	2508	19NE2	78.30	20C	2508	19CD	79.37
20C	2508	20CA	16.86	20C	2508	20N	30.40
20C	2508	21N	12.70	20C	2508	21CA	28.94
184NE1	2508	184CD1	18.63	184NE1	2508	19CG	59.96
184NE1	2508	19NE2	61.10	184NE1	2508	19CD	50.56
184NE1	2508	184CE2	16.07	184NE1	2508	20N	95.41
184NE1	2508	184CG	27.36	184CD1	2508	19CG	54.74
184CD1	2508	19NE2	67.50	184CD1	2508	19CD	52.79
184CD1	2508	184CE2	28.00	184CD1	2508	20CA	98.14
184CD1	2508	20N	81.88	184CD1	2508	184CG	15.43
19CG	2508	19NE2	32.50	1.9CG	2508	1.9CD	19.73
19CG	2508	184CE2	76.01	19CG	2508	20CA	58.57
19CG	2508	20N	42.57	19CG	2508	184CG	68.67
19CG	2508	21N	75.23	19C <b>G</b>	2508	21CA	84.51
19NE2	2508	19CD	17.06	19NE2	2508	184CE2	74.74
19NE2	2508	20CA	80.50	19 <b>NE</b> 2	2508	20N	68.75
19NE2	2508	184CG	82.83	19NE2	2508	21N	83.53
19NE2	2508	21CA	83.71	19CD	2508	184CE2	66.00
19CD	2508	20CA	76.51	19CD	2508	20N	61.57
19CD	2508	184CG	68.20	19CD	2508	21N	87.76
19CD	2508	21CA	92.52	184CE2	2508	184CG	27.48
20CA	2508	2011	17.27	20CA	2508	21N	28.61
20CA	2508	21CA	45.46	20 <b>N</b>	2508	184CG	88.95

			T.	ABLE XIII			
20N	2508	21N	43.10	20N	2508	21CA	58.85
21N	2508	21CA	16.97	200	25C9	19NE2	82.43
200	25C9	19CD	79.79	200	25C9	19CG	60.88
200	25C9	20C	3.58	200	25C9	220	59.91
200	25C9	184CD1	98.95	200	25C9	190E1	91.12
200	25C9	22N	39.58	200	25C9	21CA	31.39
200	25C9	210E1	65.02	19NE2	25C9	19CD	19.36
19NE2	25C9	19CG	36.33	19NE2	25C9	184NE1	69.05
19NE2	25C9	20C	86.00	19NE2	25C9	220	36.71
19NE2	25C9	184CD1	71.62	19NE2	25C9	190E1	26.23
19NE2	25C9	22N	69.60	19NE2	25C9	184CE2	82.17
19NE2	25C9	21CA	96.36	19CD	25C9	19CG	21.85
19CD	25C9	184NE1	54.74	19CD	25C9	20C	83.26
19CD	25C9	220	50.79	19CD	25C9	184CD1	53.82
19CD	25C9	190E1	12.53	19CD	25 <b>C</b> 9	22N	79.86
19CD	25C9	184CE2	69.35	19CG	25C9	184NE1	61.67
19CG	25C9	20C	64.10	19CG	25C9	220	53.24
19CG	25 <b>C</b> 9	184CD1	53.39	19CG	25C9	190E1	30.83
19CG	25 <b>C</b> 9	22N	72.73	19CG	25C9	184CE2	76.64
19CG	25C9	21CA	87.99	184NE1	25C9	184CD1	16.96
184NE1	25C9	190E1	43.53	184NE1	25C9	184CE2	15.12
20C	25C9	220	62.84	20C	25C9	190E1	94.48
20C	25C9	22N	40.82	20C	25C9	21CA	29.72
20C	25C9	210E1	62.22	220	25C9	190E1	61.50
220	25C9	22N	33.67	220	25C9	21CA	61.98
220	25C9	210E1	90.36	184CD1	25C9	190E1	45.59
184CD1	25C9	184CE2	27.01	190E1	25 <b>C</b> 9	22N	92.00
190E1	25C9	184CE2	57.64	22N	25C9	21CA	28.69
22N	25C9	210E1	58.53	21CA	25C9	210E1	36.88
200	25010	220	85.43	200	25010	19CD	97.70
200	25010	19CG	73.17	200	25010	22N	56.64
200	25010	20C	5.18	200	25010	22C	86.37
200	25010	21CA	39.27	200	25010	21C	51.49
200	25010	21N	20.36	200	25010	22CA	70.79
200	25010	19CB	67.40	200	25010	20 <b>N</b>	28.18
200	25010	20CA	10.76	200	25010	210E1	68.78
19NE2	25010	220	52.26	19NE2	25010	19CD	20.65
19NE2	25010	19CG	42.35	19NE2	25010	22N	96.61

•			T	ABLE XIII		•	•
19NE	2 25010	22C	66.07	19NE2		190E1	22 01
19NE	2 25010	22CA	82.67	19NE2			22.91 43.12
19NE2	25010	20N	84.11	19 <b>NE</b> 2	•		64.93
19NE2	25010	184NE1	62.68	19NE2			53.12
220	25010	19CD	65.17	220	25010		68.00
220	25010	22N	45.38	220	25010		81.85
220	25010	22C	15.92	220	25010	_	79.66
220	25010	21C	61.19	220	25010		79.35
220	25010	190E1	72.72	220	25010	22CA	30.41
220	25010	19CB	55.31	220	25010	·20N	77.38
220	25010	23N	24.55	220	25010	20CA	82.51
220	25010	23CA	30.74	19CD	25010	19CG	24.88
19CD	25010	20C	99.46	19CD	25010	22C	80.69
19CD	25010	190E1	10.13	19CD	25010	22CA	93.93
19CD	25010	19CB	31.83	19CD	25010	20N	69.77
19CD	25010	23N	82.79	19CD	25010	20CA	87.09
19CD	25010	184NE1	49.90	19CD	25010	23CA	72.96
19CG	25010	22N	93.59	19CG	25010	20C	75.37
19CG	25010	22C	83.51	19CG	25010	21N	89.04
19CG	25010	190E1	32.51	19CG	25010	22CA	89.71
19CG	25010	19CB	14.55	19CG	25010	20N	45.05
19CG	25010	23N	91.20	19CG	25010	20CA	62.46
19CG	25010	184NE1	58.10	19CG	25010	23CA	87.28
22N	25010	20C	51.51	22N	25010	22C	36.02
22N	25010	21CA	34.76	22N	25010	21C	15.90
22N	25010	21N	40.79	22N	25010	22CA	16.92
22N	25010	19CB	79.30	2 <b>2N</b>	25010	20N	69.27
22N	25010	23N	46.59	22N	25010	20CA	61.11
22N	25010	210E1	66.06	22N	25010	23CA	63.27
20C	25010	22C	81.89	20C	25010	21CA	35.03
20C	25010	21C	46.40	20C	25010	21N	15.83
20C	25010	22CA	65.90	20C	25010	19CB	68.42
20C	25010	20N	30.90	20C	25010	23N	95.35
20C	25010	29CA	14.07	200	25010	210E1	66.77
22C	25010	21CA	70.57	22C	25010	21C	50.65
22C	25010	21N	75.18	22C	25010	190E1	87.67
22C	25010	22CA	19.15	22C	25010	19CB	70.26
22C	25010	20N	85.66	220	25010	23N	14.18

			TA	BLE XIII			
22C	25010	20CA	86.38	22C	25010	210E1	94.63
22C	25010	23CA	28.51	21CA	25010	21C	20.23
21CA	25010	21N	19.26	21CA	25010	22CA	51.43
21CA	25010	19CB	95.98	21CA	25010	20N	65.09
21CA	25010	23N	79.47	21CA	25010	20CA	49.07
21CA	25010	210E1	39.22	21CA	25010	23CA	96.32
21C	25010	21N	32.16	21C	25010	22CA	31.67
21C	25010	19CB	90.80	21C	25010	20N	71.25
21C	25010	23N	59.25	21C	25010	20CA	58.84
21C	25010	210E1	50.69	21C	25010	23CA	76.11
21N	25010	22CA	57.03	21N	25010	19CB	80.04
21N	25010	20 <b>N</b>	45.98	21N	25010	23N	87.15
21N	25010	20CA	29.82	21N	25010	210E1	54.21
190El	25010	19CB	41.46	190E1	25010	20N	77.43
190E1	25010	23N	87.76	190E1	25010	20CA	94.86
190E1	25010	184NE1	41.58	190E1	25010	23CA	75.81
22CA	25010	19CB	75.23	22CA	25010	20N	77.28
22CA	25010	23N	30.13	22CA	25010	20CA	73.19
22CA	25010	210E1	78.69	22CA	25010	23CA	46.56
19CB	25010	20N	41.01	19CB	25010	23N	79.42
19CB	25010	20CA	57.32	19CB	25010	184NE1	72.38
19CB	25010	23CA	78.47	20 <b>N</b>	25010	23N	99.51
20N	25010	20CA	17.43	20N	25010	184NE1	88.50
20N	25010	210E1	96.89	23N	25010	210E1	95.90
23N	25010	23CA	16.87	20CA	25010	210E1	79.48
19NE2	25C11	19CD	13.29	19NE2	25C11	184NE1	59.74
19NE2	25C11	220	32.16	19CD	25C11	184NE1	46.55
19CD	25C11	220	44.12	184NE1	25C11	220	89.70
19NE2	25C12	220	39.61	19NE2	25C12	23CA	60.70
19NE2	25C12	22C	55.91	19NE2	25C12	23N	65.25
19NE2	25C12	224OH2	93.02	19NE2	25C12	22 <b>N</b>	67.68
1.9NE2	25C12	19CD	8.19	19NE2	25C12	200	56.75
220	25C12	23CA	39.57	220	25C12	22C	16.82
220	25C12	23N	31.30	220	25C12	224OH2	75.08
220	25C12	22N	34.48	220	25C12	19CD	46.01
220	25C12	200	50.49	23CA	25C12	22C	33.54
23CA	25C12	23N	19.76	23CA	25C12	2240H2	36.28
23CA	25C12	22N	62.96	23CA	25C12	19CD	68.87

			TA	ABLE XIII			
23CA	25C12	200	89.77	22C	25C12	23N	17.94
22C	25C12	2240H2	64.69	22C	25C12	22N	29.81
22C	25C12	19CD	62.67	22C	25C12		58.58
23N	25C12	224OH2	46.80	23N	25C12		44.68
23N	25C12	19CD	73.05	23N	25C12	200	76.44
224OH2	25C12	22N	89.06	22N	25C12	19CD	71.05
22N	25C12	200	37.38	19CD	25C12	200	54.60
210E1	25C13	22N	66.00	210E1	25C13	22C	95.57
210E1	25C13	220	97.18	210E1	25C13	21C	51.73
22N	25C13	22C	29.65	22N	25C13	23N	43.65
22N	25C13	220	33.57	22N	25C13	21C	15.61
22N	25C13	23CA	58.56	22C	25C13	23N	16.28
22C	25C13	220	14.98	22C	25C13	210	44.10
22C	25C13	23CA	29.00	23N	25C13	220	27.57
23N	25C13	21C	56.18	23N	25C13	23CA	17.03
220	25C13	21C	49.18	220	25C13	23CA	33.12
21C	25C13	23CA	72.30	22N	25C14	21C	21.24
22N	25C14	22C	38.14	22N	25C14	210E1	82.54
22N	25C14	23N	56.21	22N	25C14	22CA	22.02
22N	25C14	21CA	36.01	2 <b>2N</b>	25C14	210	33.22
22N	25C14	220	40.42	22N	25C14	23CA	70.62
22N	25C14	200	41.05	22N	25C14	21CB	49.85
22N	25C14	21CD	75.68	21C	25C14	22C	58.12
21C	25C14	210E1	65.17	21C	25C14	23N	74.13
21C	25C14	22CA	37.70	21.C	25C14	21CA	21.82
21C	25C14	210	17.02	21C	25C14	220	61.64
21C	25C14	23CA	90.21	21C	25C14	200	46.27
21C	25C14	21CB	30.54	21C	25C14	21CD	59.45
22C	25C14	23N	20.02	22C	25C14	22CA	22.87
22C	25C14	21CA	73.72	22C	25C14	210	63.62
22C	25C14	220	16.99	22C	25C14	23CA	32.48
22C	25C14	200	63.32	22C	25C14	21CB	87.93
210E1	25C14	21CA	46.55	210 <b>F1</b>	25C14	210	68.11
210E1	25C14	200	64.41	210E1	25C14	21CB	34.92
210E1	25C14	21CD	7.92	23N	25C14	22CA	36.43
23N	25C14	21CA	92.21	23N	25C14	210	75.11
23N	25C14	220	32.42	23N	25C14	23CA	17.87
23N	25C14	200	82.87	22CA	25C14	21CA	

		TA	ABLE XIII			
22CA	25C14 210	40.83	22CA	25C14	220	34.41
22CA	25C14 23CA	53.03	22CA	25C14	200	60.12
22CA	25C14 21CB	68.24	22CA	25C14	21CD	96.45
21CA	25C14 210	34.25	21CA	25C14	220	71.57
21CA	25C14 200	35.31	21CA	25C14	21CB	17.17
21CA	25C14 21CD	39.72	210	25C14	220	71.78
210	25C14 23CA	92.77	210	25C14	200	63.21
210	25C14 21CB	34.50	210	25C14	21CD	64.40
220	25C14 23CA	37.01	220	25C14	200	51.83
220	25C14 21CB	88.01	23CA	25C14	200	88.32
200	25C14 21CB	50.65	200	25C14	21CD	56.60
21CB	25C14 21CD	30.13	210E1	25C15	21CD	12.20
210E1	25C15 21NE2	26.50	210E1	25C15	21CA	39.20
210E1	25C15 200	65.35	21CD	25C15	21NE2	16.43
21CD	25C15 21CA	39.53	21CD	25C15	200	59.46
21NE2	25C15 21CA	53.49	21NE2	25C15	200	65.84
21CA	25C15 200	32.72	19NE2	25C16	19CD	16.53
19NE2	25C16 162ND1	88.72	19NE2	25C16	190E1	31.45
19NE2	25C16 162CE1	76.13	19NE2	25C16 1	84NE1	68.38
19NE2	25C16 184CE2	84.59	19NE2	25C16	25 <b>S</b> G	70.09
19NE2	25C16 23CA	54.01	19NE2	25C16	220	29.07
19CD	25C16 162ND1	77.07	19CD	25C16	190E1	16.88
19CD	25C16 162CE1	62.31	19CD	25C16 1	84NE1	52.74
19CD	25C16 184CZ2	85.13	19CD	25C16 1	84CE2	69.00
19CD	25C16 25SG	69.54	19CD	25C16	23CA	69.33
19CD	25016 220	45.46	162ND1	25C16	190E1	60.33
162ND1	25C16 162CE1	17.92	162ND1	25C16 1	84NE1	63.79
162ND1	25C16 184CZ2	60.49	162ND1	25C16 1	84CE2	63.53
162ND1	25C16 25SG	43.67	190E1	25C16 1	62CE1	45.48
190E1	25C16 184NE1	45.86	190E1	25C16 1	84CZ2	75.36
	25C16 184CE2	61.18	190E1	25C16	25 <b>S</b> G	59.93
190E1	25C16 23CA	78.49	190 <b>E1</b>	25C16	220	60.19
162CE1	25C16 184NE1	47.41	162CE1	25016 1	84CZ2	52.59
162CE1	25C16 184CE2	50.53	162CE1	25C16	25 <b>S</b> G	51.17
184NE1	25C16 184CZ2	32.85	184NE1	25C16 1	84CE2	16.26
1.84NE1	25C16 25SG	93.75	184NE1	25C16	220	94.69
184CZ2	25C16 184CE2	17.00	25 <i>S</i> G	25C15	2 <b>3ÇA</b>	63.45
25SG	25C16 22O	79.20	23CA	25C16	220	32.47

		T.	ABLE XIII		
19NE2		87.64	19NE2	25017 162CE1	97.12
19NE2	25017 190E1	40.11	19NE2	25017 19CD	22.12
19NE2	25017 184CD1	77.66	19NE2	25017 25CB	62.38
19NE2	25017 25 <i>S</i> G	74.14	19NE2	25017 19CG	26.63
184NE1	25017 162CE1	65.10	184NE1	25017 162ND1	86.13
184NE1	25017 190E1	60.63	184NE1	25017 19CD	67.63
184NE1	25017 184CZ2	42.58	184NE1	25017 184CE2	20.97
184NE1	25017 162NE2	55.54	184NE1	25017 162CG	83.50
184NE1	25017 184CD1	10.23	184NE1	25017 25CB	94.14
184NE1	25017 19CG	61.07	184NE1	25017 162CD2	67.31
184NE1	25017 184CH2	48.57	162CE1	25017 162ND1	23.08
162CE1	25017 190E1	59.61	162CE1	25017 19CD	80.14
162CE1	25017 184CZ2	69.34	162CE1	25017 184CE2	67.58
162CE1	25017 162NE2	14.63	162CE1	25017 162CG	29.19
162CE1	25017 184CD1	69.18	162CE1	25017 25CB	46.12
162CE1	25017 25SG	57.25	162CE1	25017 19CG	86.83
162CE1	25017 162CD2	22.83	162CE1	25017 184CH2	73.20
162ND1	25017 190E1	77.34	162ND1	25017 19CD	96.95
162ND1	25017 184CZ2	77.78	162ND1	25017 184CE2	83.59
162ND1	25017 162NE2	30.71	162ND1	25017 162CG	13.98
162ND1	25017 184CD1	91.47	162ND1	25017 25CB	47.42
162ND1	25017 25SG	46.98	162ND1	25017 162CD2	23.75
162ND1	25017 184CH2	79.38	190E1	25017 19CD	20.54
190E1	25017 184CZ2	99.66	190E1	25017 184CE2	80.49
190E1	25017 162NE2	66.69	190E1	25017 162CG	88.06
190E1	25017 184CD1	53.52	190E1	25017 25CB	44.71
190E1	25017 25SG	67.03	190E1	25017 19CG	28.75
190E1	25017 162CD2	81.23	19CD	25017 184CE2	88.60
19CD	25017 162NE2	86.77	19CD	25017 184CD1	58.23
	25017 25CB		19CD	25017 25SG	76.46
19CD	25017 19CG		184CZ2	25017 184CE2	21.82
	25017 162NE2		184CZ2	25017 162CG	66.69
	25017 184CD1	52.41	184CZ2	25017 162CD2	54.17
		6.39	184CE2	25017 162NE2	54.32
184CE2	25017 162CG	76.31	184CE2	25017 184CD1	30.61
	25017 19CG		184CE2	25017 162CD2	60.79
	25017 184CH2		162NE2	25017 162CG	29.72
162NE2	25017 184CD1	61.67	162NE2	25017 25CB	60.57

		Т	ABLE XIII		
162NE2	25017 25SG	71.37	162NE2	25017 19CG	90.84
162NE2	25017 162CD2	15.82	162NE2	25017 184CH2	58.64
162CG	25017 184CD1	90.78	162CG	25017 25CB	61.39
162CG	25017 25SG	59.43	162CG	25017 162CD2	16.23
162CG	25017 184CH2	67.39	184CD1	25017 25CB	91.37
184CD1	25017 19CG	51.04	184CD1	25017 162CD2	74.86
184CD1	25017 184CH2	58.22	25CB.	25017 25SG	23.19
25CB	25017 19CG	69.56	25 <b>C</b> B	25017 162CD2	65.94
25SG	25017 19CG	88.74	25 <i>S</i> G	25017 162CD2	70.29
162CD2	25017 184CH2	56.25	19NE2	25N18 25SG	74.49
19NE2	25N18 162ND1	81.92	19NE2	25N18 23CA	57.55
19NE2	25N18 224OH2	93.53	19NE2	25N18 162CE1	67.31
19NE2	25N18 19CD	12.59	19NE2	25N18 25CB	55.64
19NE2	25N18 19OE1	26.71	25SG	25N18 162ND1	48.92
25 <i>S</i> G	25N18 1610	67.41	25SG	25N18 23CA	74.24
25 <b>S</b> G	25N18 224OH2	87.02	25 <i>S</i> G	25N18 162CE1	53.82
25SG	25N18 19CD	69.72	25SG	25N18 25CB	21.75
25SG	25N18 19OE1	60.33	162ND1	25N18 1610	62.58
162ND1	25N18 162CE1	16.19	162ND1	25N18 19CD	69.87
162ND1	25N18 25CB	43.26	162ND1	25N18 190E1	55.30
1610	25N18 224OH2	98.16	1610	25N18 162CE1	78.73
1610	25N18 25CB	84.03	23CA	25N18 224OH2	36.29
23CA	25N18 19CD	68.30	23CA	25N18 25CB	74.55
23CA	25N18 19OE1	77.72	2240H2	25N18 25CB	98.57
162CE1	25N18 19CD	54.91	162CE1	25N18 25CB	40.63
162CE1	25N18 19OE1	40.66	19CD	25N18 25CB	48.94
19CD	25N18 190E1	14.63	25CB	25N18 19OE1	38.63
25 <i>S</i> G	25N19 162ND1	68.67	25\$G	25N19 1610	97.88
25SG	25N19 25CB	26.20	25 <b>S</b> G	25N19 162CE1	69.25
25 <i>S</i> G	25N19 162CA	69.30	25SG	25N19 162CG	76.89
25SG	25N19 19NE2	81.47	25 <b>S</b> G	25N19 162CB	81.47
25SG	25N19 161C	91.20	25SG	25N19 23CA	80.48
25 <b>S</b> G	25N19 162N	79.54	25 <i>S</i> G	25N19 190E1	67.38
25 <i>S</i> G	25N19 224OH2	96.97	1.62ND1	25N19 1610	84.02
162ND1	25N19 25CB	55.97	162ND1	25N19 162CE1	
	25N19 162CA	46.46	162ND1	25N19 162CG	15.15
162ND1	25N19 19NE2	86.65	162ND1	25N19 162CB	
162ND1	25N19 161C	76.02	162ND1	25N19 162N	

		1	ABLE XII	I	
162ND1		59.63		25N19 162CA	41.35
1610	25N19 162CG	69.81	1610	25N19 162CB	49.97
1610	25N19 161C	8.90	1610	25N19 162N	24.76
25CB	25N19 162CE1	49.59	25CB	25N19 162CA	78.03
25CB	25N19 162CG	68.95	25CB	25N19 19NE2	60.40
25CB	25N19 162CB	81.67	25CB	25N19 23CA	80.10
25CB	25N19 162N	92.77	25CB	25N19 19OE1	41.65
162CE1		63.58	162CE1		31.41
162CE1	25N19 19NE2	69.42	162CE1		51.18
162CE1	25N19 161C	93.28	162CE1		79.16
162CE1	25N19 19OE1	42.84	162CA	25N19 162CG	35.79
162CA	25N19 162CB	20.40	162CA	25N19 161C	32.47
162CA	25N19 162N	16.68	162CG	25N19 162CB	19.86
162CG	25N19 161C	62.27	162CG	25N19 162N	49.28
162CG	25N19 19OE1	74.24	19NE2	25N19 23CA	51.93
19NE2	25N19 19OE1	27.73	19NE2	25N19 224OH2	83.06
162CB	25N19 161C	42.44	162CB	25 <b>N1</b> 9 162N	30.28
162CB	25N19 19OE1	93.88	161C	25N19 162N	15.85
23CA	25N19 19OE1	74.91	23CA	25N19 224OH2	33.48
19NE2	25N20 184NE1	66.71	19NE2	25N20 19CD	16.40
19NE2	25N20 200	61.92	19NE2	25N20 184CE2	83.41
19NE2	25N20 184CZ2	92.74	19NE2	25N20 184CD1	65.86
19NE2	25N20 19OE1	25.29	19NE2	25N20 19CG	28.13
184NE1	25N20 19CD	51.81	184NE1	25N20 200	92.93
184NE1	25N20 184CE2	17.64	184NE1	25N20 184CZ2	33.11
184NE1	25N20 184CD1	14.57	184NE1	25N20 19OE1	41.63
184NE1	25N20 19CG	54.80	19CD	25N20 200	61.53
19CD	25N20 184CE2	69.14	19CD	25N20 184CZ2	80.93
19CD	25N20 184CD1	49.60	19CD	25N20 19OE1	13.44
19CD	25N20 19CG	17.37	200	25N20 184CD1	79.82
200	25N20 19OE1	74.00	200	25N20 19CG	46.41
134CE2	25N20 184CZ2	17.72	184CE2	25N20 184CD1	27.30
184CE2	25N20 19OE1	58.12	184CE2	25N20 19CG	72.23
184CZ2	25N20 184CD1	44.82	184CZ2	25N20 19OE1	
184CZ2	25N20 19CG	87.39	184CD1		43.08
184CD1	25N20 19CG	47.27		25N20 19CG	27.95
25SG	25C21 25CB	34.16	25 <b>5</b> G	25C21 25N	59.13
25 <b>S</b> G	25C21 162ND1	63.16	25 <b>S</b> G	25C21 230	95.53

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			TA	ABLE XIII		
25SG	25C21	. 25CA	41.14	25SG	25C21 161C	90.20
25SG	25C21	162CE1	64.32	25 <b>S</b> G	25C21 24N	94.43
25SG	25C21	190E1	78.34	25 <b>S</b> G	25C21 162C	A 58.84
25SG	25C21	19CD	91.01	25SG	25C21 26N	32.55
25SG	25C21	25C	28.93	25SG	25C21 24C	64.37
25 <b>S</b> G	25C21	162CG	63.81	25SG	25C21 163N	30.18
25CB	25C21	25N	41.43	25CB	25C21 162N	D1 57.90
25CB	25C21	23C	92.08	25CB	25C21 230	94.39
25CB	25C21	25CA	20.31	25CB	25C21 19N	E2 70.68
25CB	25C21	162CE1	48.50	25CB	25C21 24N	77.34
25CB	25C21	190E1	44.80	25CB	25C21 162C	A 77.90
25CB	25C21	19CD	56.86	25CB	25C21 26N	42.39
25CB	25C21	25C	26.69	25CB	25C21 24C	49.86
25CB	25C21	162CG	64.53	25CB	25C21 163N	55.35
25N	25C21	162ND1	98.16	25N	25C21 23C	A 69.54
25N	25C21	23C	50.81	25N	25C21 230	54.24
25N	25C21	25CA	21.67	25N	25C21 19N	E2 62.10
25N	25C21	162CE1	85.44	25N	25C21 24N	37.13
25N	25C21	190E1	55.93	25 <b>N</b>	25C21 19C	56.26
25N	25C21	26N	35.66	25N	25C21 25C	30.32
25N	25C21	24C	8.74	25N	25C21 163N	88.85
162ND1	25C21	25 <b>CA</b>	78.14	162ND1	25C21 19N	84.09
162ND1	25C21	1610	65.36	162ND1	25C21 162C1	E1 16.67
162ND1	25C21	190E1	59.98	162ND1	25C21 162C	39.67
162ND1	25C21	19CD	72.99	162ND1	25C21 26N	92.87
162ND1	25C21	25C	80.82	162ND1	25C21 162C	9.83
162ND1	25C21	163N	47.12	23CA	25C21 23C	22.59
23CA	25C21	230	35.84	23CA	25C21 25C2	90.84
23CA	25C21	19NE2	61.29	23CA	25C21 224OF	38.65
23CA	25C21	24N	32.44	23CA	25C21 19OF	87.52
23CA	25C21	19CD	73.32	23CA	25C21 26N	93.91
23CA	25C21	25C	97.86	23CA	25C21 24C	62.65
23C	25C21	230	18.06	23C	25C21 25CF	72.44
23C	25C21	19NE2	67.14	23C	25C21 224OF	12 52.52
23C	25C21	24N	16.23	23C	25C21 19OF	85.86
23C	25C21	19CD	74.76	23C	25C21 26N	71.38
23C	25C21	25C	76.60	23C	25C21 24C	42.81
230	25C21	25 <b>CA</b>	74.11	230	25C21 19NE	2 84.69

		Т	ABLE XIII		
230	25C21 224OH2	51.69	230	25C21 24N	29.95
230	25C21 19CD	90.88	230	25C21 26N	63.28
230	25C21 25C	72.90	230	25C21 24C	45.50
25CA	25C21 19NE2	68.27	25CA	25C21 162CE1	67.59
25CA	25C21 24N	58.61	25CA	25C21 19OE1	50.41
25CA	25C21 162CA	95.22	25CA	25C21 19CD	57.40
25CA	25C21 26N	30.73	25CA	25C21 25C	16.57
25CA	25C21 24C	29.77	25CA	25C21 162CG	84.82
25CA	25C21 163N	69.11	19NE2	25C21 162CE1	68.48
19NE2	25C21 224OH2	93.44	19NE2	25C21 24N	55.94
19NE2	25C21 19OE1	29.51	19NE2	25C21 19CD	14.63
19NE2	25C21 26N	96.37	19NE2	25C21 25C	84.76
19NE2	25C21 24C	64.60	19NE2	25C21 162CG	93.25
1610	25C21 162CE1	81.61	1610	25C21 162CA	35.10
1610	25C21 162CG	55.78	1610	25C21 163N	60.78
162CE1	25C21 19OE1	43.32	162CE1	25C21 162CA	55.76
162CE1	25C21 19CD	56.63	162CE1	25C21 26N	88.80
162CE1	25C21 25C	74.35	162CE1	25C21 24C	94.10
162CE1	25C21 162CG	26.49	162CE1	25C21 163N	57.15
224OH2	25C21 24N	67.70	2240H2	25C21 24C	94.80
24N	25C21 19OE1	70.64	24N	25C21 19CD	60.95
24N	25C21 26N	64.52	24N	25C21 25C	65.81
24N	25C21 24C	30.33	190E1	25C21 162CA	98.48
190E1	25C21 19CD	15.08	190E1	25C21 26N	81.12
190E1	25C21 25C	66.02	1.90E1	25C21 24C	62.63
190E1	25C21 162CG	69.79	190E1	25C21 163N	90.46
162CA	25C21 26N	39.91	162CA	25C21 25C	87.29
162CA	25C21 162CG	30.51	1.62CA	25C21 163N	28.80
19CD	25C21 26N	87.36	19CD	25C21 25C	73.91
19CD	25C21 24C	60.95	19CD	25C21 162CG	82.63
26N	25C21 25C	15.95	26N	25C21 24C	37.06
26N	25C21 162CG	95.38	2 <b>6N</b>	25C21 163N	61.27
25C	25C21 24C	35.55	25C	25C21 162CG	85.16
25C	25C21 163N	58.98	24C	25C21 163N	94.51
162CG	25C21 163N	42.83	25SG	25022 25CB	38.49
25SG	25022 25N	68.14	25 <b>SG</b>	25022 25CA	51.61
25SG	25022 230	92.70	25 <b>S</b> G	25022 190E1	92.24
25SG	25022 162ND1	54.00	25SG	25022 24C	76.75

			TA	ABLE XIII			
25 <i>S</i> G	25022	162CE1	62.63	25SG	2502	2 24CA	92.42
25SG	25022	25C	41.17	25SG	2502	2 26N	39.32
19NE2	25022	25CB	90.81	19NE2	2502	2 23CA	83.72
19NE2	25022	25N	82.91	19NE2	25022	2 23C	90.92
19NE2	25022	25CA	85.84	19NE2	25022		75.03
19NE2	25022	19CD	18.29	19NE2	25022		36.86
19NE2	25022	220	40.88	19NE2	25022		93.79
19NE2	25022	24C	82.04	19NE2	25022	23N	71.12
19NE2	25022	162CE1	77.66	19NE2	25022	24CA	80.86
19NE2	25022	22C	54.65	19NE2	25022	25C	99.55
25CB	25022	25N	47.57	25CB	25022	25CA	24.14
25CB	25022	24N	93.92	25CB	25022	19CD	72.69
25CB	25022	190E1	55.94	25CB	25022	162ND1	53.00
25CB	25022	24C	58.84	25CB	25022	162CE1	47.93
25CB	25022	24CA	78.52	25CB	25022	25C	26.37
25CB	25022	26N	39.92	23CA	25022	25N	86.84
23CA	25022	23C	27.45	23CA	25022	24N	40.92
23CA	25022	230	40.51	23CA	25022	19CD	98.21
23CA	25022	220	43.20	23CA	25022	24C	75.56
23CA	25022	23N	12.75	23CA	25022	2240H2	38.92
23CA	25022	24CA	55.88	23CA	25022	22C	29.08
25N	25022	23C	61.42	25 <b>N</b>	25022	25CA	23.44
25N	25022	24N	46.54	25 <b>N</b>	25022	230	61.31
25N	25022	19CD	72.72	25N	25022	190E1	69.34
25N	25022	220	78.77	25N	25022	24C	11.28
25N	25022	23N	87.24	25N	25022	162CE1	91.85
25N	25022	24CA	30.96	25N	25022	22C	83.18
25N	25022	25C	27.52	25N	25022	26N	31.88
23C	25022	25CA	84.51	23C	25022	24N	21.22
23C	25022	230	19.47	23C	25022	19CD	98.72
23C	25022	220	52.92	23C	25022	24C	50.34
23C	25022	23N	34.26	23C	25022	2240H2	55.48
23C		24CA	31.37	23C	25022	22C	43.43
23C	25022	25C	83.00	23C	25022	26N	74.42
25CA	25022	24N	69.81	25CA	25022	230	81.49
25CA	25022	19CD	70.36	25CA	25022	190E1	59.53
25CA	25022	220	96.17	25CA	25022	162ND1	76.96
25CA	25022	24C	34.73	25CA	25022	162CE1	69.68

			т	ABLE XIII			
25CA	25022	24CA	54.41	25CA	25022	25C	13.77
25CA	25022	26N	28.59	24N	25022		35.89
24N	25022	19CD	79.36		25022		90.36
24N	25022	220	44.09	24N	25022		35.42
24N	25022	23N	40.92	24N	25022	_	75.53
24N	25022	24CA	16.33	24N	25022		41.01
24N	25022	25C	72.48	24N	25022		69.02
230	25022	220	72.38	230	25022		51.99
230	25022	23N	50.70	230	25022		53.17
230	25022	24CA	37.85	230	25022		62.29
230	25022	25C	75.63	230	25022	_	63.41
19CD	25022	190E1	19.08	19CD	25022	220	55.19
19CD	25022	162ND1	80.93	19CD	25022	24C	75.07
19CD	25022	23N	86.19	19CD	25022		63.77
19CD	25022	24CA	80.05	19CD	25022	22C	69.54
19CD	25022	25C	84.09	19CD	25022	26N	98.42
190E1	25022	220	73.71	190E1	25022	162ND1	64.15
190E1	25022	24C	75.30	190E1	25022	162CE1	46.77
190E1	25022	24CA	86.24	190E1	25022	22C	88.09
190E1	25022	25C	72.43	190E1	25022	26N	87.99
220	25022	24C	70.73	220	25022	23N	31.00
220	25022	224OH2	78.01	220	25022	24CA	57.51
220	25022	22C	14.38	162ND1	25022	162CE1	17.38
162ND1	25022	25C	77.53	162ND1	25022	26N	86.27
24C	25022	23N	76.28	24C	25022	24CA	19.68
24C	25022	22C	73.43	24C	25022	25C	37.56
24C	25022	26N	38.26	23 <b>N</b>	25022	2240H2	48.26
23N	25022	24CA	57.15	23N	25022	22C	16.66
162CE1	25022	25C	74.28	162CE1	25022	26N	86.57
2240H2	25022	24CA	86.57	224OH2	25022	22C	63.94
24CA	25022	22C	56.72	24CA	25022	25C	56.18
24CA	25022	26N	53.14	25C	25022	26N	15.75
610D1	25C23	590	92.71	610D1	25C23	2640H2	51.41
610D1	25C23	61CG	13.18	610D1	25C23	610D2	26.94
610D1	25C23	59C	87.40	590	25C23	67CD2	91.32
590	25C23	2640H2	79.96	590	25C23	61CG	79.93
590	25C23	61002	71.08	59C	25C23	59C	5.39
67CE2	25C23	67CD2	18.11	67CD2	25C23	59C	92.59

			TA	BLE XIII			
2640H2	25C23	61CG	46.02	264OH2	25C23	610D2	33.75
2640H2	25C23	59C	77.27	61CG	25C23	610D2	15.44
61CG	25C23	59C	74.68	610D2	25C23	59C	66.25
590	25C24	60ND2	58.48	590	25C24	60CA	44.55
590	25C24	59C	9.80	590	25C24	61CG	88.32
590	25C24	60C	54.30	590	25C24	61N	71.19
590	25C24	60N	26.62	590	25C24	700D1	52.26
590	25C24	610D2	75.43	590	25C24	60CG	60.01
590	25C24	60CB	55.47	590	25C24	67N	96.69
610D1	25C24	60CA	80.24	610D1	25C24	59¢	97.84
610D1	25C24	61CG	15.04	610D1	25C24	66CA	87.60
610D1	25C24	60C	61.23	610D1	25C24	61N	50.12
610D1	25C24	60N	89.60	610D1	25C24	610D2	27.04
610D1	25C24	60CB	87.33	610D1	25C24	650	55.48
67CD2	25C24	60ND2	70.35	67CD2	25C24	67CE2	21.12
67CD2	25C24	66CA	60.05	67CD2	25C24	700D1	65.25
67CD2	25C24	60CG	77.71	67CD2	25C24	60CB	94.80
67CD2	25C24	6 <b>7N</b>	38.69	67CD2	25C24	66C	42.86
67CD2	25C24	650	91.74	67CD2	25C24	67CG	7.62
60ND2	25C24	67CE2	91.22	60ND2	25C24	60CA	41.72
60ND2	25C24	59C	52.47	60ND2	25C24	66CA	51.12
60ND2	25C24	60C	59.58	60ND2	25C24	61N	66.53
60ND2	25C24	60 <b>N</b>	44.60	60ND2	25C24	700D1	41.24
60ND2	25C24	60CG	12.36	60ND2	25C24	60CB	29.62
60ND2	25C24	67N	38.69	60ND2	25C24	66C	49.79
60ND2	25C24	650	68.79	60ND2	25C24	67CG	64.10
67CE2	25C24	66CA	70.56	67CE2	25C24	700D1	82.90
67CE2	25C24	60CG	97.67	67CE2	25C24	67 <b>N</b>	56.69
67CE2	25C24	66C	55.71	67CE2	25C24	650	96.69
67CE2	25C24	67CG	28.20	60CA	25C24	59C	34.75
60CA	25C24	61CG	65.91	60CA	25C24	66CA	65.63
60CA	25C24	60C	19.12	60CA	25C24	61N	32.35
60CA	25024	60N	17.94	60CA	25C24	700D1	70.97
60CA	25C24	610D2	62.09	60CA	25C24	60CG	32.72
60CA	25C24	60CB	16.35	50CA	25C24	67N	73.13
60CA	25C24	66C	76.33	60CA	25C24	650	56.57
59 <b>C</b>	25C24	61CG	83.18	59C	25C24	66CA	96.04
59C	25C24	60C	45.42	59C	25C24	61N	62.28

			TA	BLE XIII			
59C	25C24	60N	16.82	59C	25C24	700D1	54.54
59C	25C24	610D2	71.72	59C	25C24	60CG	52.21
59C	25C24	60CB	46.02	59C	25C24	67N	91.16
59C	25C24	650	91.18	61CG	25C24	66CA	85.83
61CG	25C24	60C	46.79	61CG	25C24	61N	37.73
61CG	25C24	60N	74.56	61CG	25C24	610D2	15.30
61CG	25C24	60CG	92.19	61CG	25C24	60CB	74.64
61CG	25C24	650	52.44	66CA	25C24	60C	69.53
66CA	25C24	61N	60.35	66CA	25C24	60N	81.05
66CA	25C24	700D1	85.92	66CA	25C24	610D2	97.05
66CA	25C24	60CG	45.49	66CA	25C24	60CB	50.81
66CA	25C24	67N	29.45	66CA	25C24	66C	17.50
66CA	25C24	650	33.39	66CA	25C24	67CG	60.31
60C	25C24	61N	16.90	60C	25C24	60N	31.03
60C	25C24	700D1	89.71	60C	25C24	610D2	43.83
60C	25C24	60CG	49.02	60C	25C24	60CB	30.73
60C	25C24	67 <b>N</b>	85.40	60C	25C24	66C	84.14
60C	25C24	650	49.16	61N	25C24	60N	47.32
61N	25C24	610D2	41.00	61N	25C24	60CG	54.47
61N	25C24	60CB	37.23	61N	25C24	67N	82.57
61N	25C24	66C	77.02	61 <b>N</b>	25C24	650	34.06
60N	25C24	700D1	61.08	60 <b>N</b>	25C24	610D2	66.32
60N	25C24	60CG	40.42	60N	25C24	60CB	30.28
60N	25C24	67N	81.86	60N	25C24	66C	88.80
60N	25C24	650	74.46	700D1	25C24	60CG	53.02
700D1	25C24	60CB	66.53	70001	25C24	67N	60.32
700D1	25C24	6 SC	75.88	700D1	25C24	67CG	57.76
610D2	25C24	60CG	92.66	610D2	25C24	60CB	74.33
610D2	25C24	650	64.27	60C <b>G</b>	25C24	60CB	18.33
60CG	25C24	67 <b>N</b>	41.67	60 <b>CG</b>	25C24	66C	48.95
60CG	25C24	650	57.71	60CG	25C24	67CG	72.33
60CB	25C24	67 <b>N</b>	5€.95	60CB	25C24	66C	60.17
60CB	25C24	650	49.72	60CB	25C24	67CG	90.00
67N	25C24	66C	15.58	67N	25C24	650	62.20
67N	25C24	67CG	35.72	66C	25C24	650	50.68
66C	25C24	67CG	42.81	650	25C24	67CG	93.15
66CA	25C25	67CD2	74.47	66CA	25C25	67CE2	88.53
66CA	25C25	60ND2	59.31	6ECA	25C25	66N	21.62

			TA	BLE XIII			
66CA	25C25	650	44.29	66CA	25C25	66C	21.53
66CA	25C25	65C	36.30	66CA	25C25	60CA	74.60
66CA	25C25	61N	73.02	66CA	25C25	67N	33.98
66CA	25C25	60C	79.39	66CA	25C25	60CG	50.06
66CA	25C25	660	28.68	66CA	25C25	60CB	56.47
66CA	25C25	61CB	88.35	66CA	25C25	67CG	69.03
66CA	25C25	67CZ	89.74	610D1	25C25	66N	99.44
610D1	25C25	650	70.37	610D1	25C25	65C	80.66
610D1	25C25	60CA	81.54	610D1	25C25	61CG	15.19
610D1	25C25	61N	55.29	610D1	25C25	590	88.76
610D1	25C25	60C	62.96	610D1	25C25	60CB	93.30
610D1	25C25	61CB	28.06	610D1	25C25	610D2	23.60
67CD2	25C25	67CE2	22.80	67CD2	25C25	60ND2	71.72
67CD2	25C25	66N	86.08	67CD2	25C25	66C	52.97
67CD2	25C25	590	98.99	67CD2	25C25	67N	45.03
67CD2	25C25	60CG	81.76	67CD2	25C25	660	50.77
67CD2	25C25	67CG	6.70	67CD2	25C25	67CZ	27.35
67CE2	25C25	60ND2	94.29	67CE2	25C25	6 <b>6N</b>	93.17
67CE2	25C25	66C	67.80	67CE2	25C25	67N	64.98
67CE2	25C25	660	60.61	67CE2	25C25	67CG	29.23
67CE2	25C25	67CZ	4.94	60ND2	25C25	66N	80.00
60ND2	25C25	650	81.55	60 <b>N</b> D2	25C25	66C	58.26
60ND2	25C25	65C	87.34	60ND2	25C25	60CA	40.17
60ND2	25C25	61N	69.84	60ND2	25C25	590	49.72
60ND2	25C25	67 <b>N</b>	43.92	60ND2	25C25	60C	58.97
60ND2	25C25	60CG	15.02	60MD2	25C25	660	71.48
60ND2	25C25	60CB	31.11	60ND2	25C25	67CG	65.78
60ND2	25C25	610D2	99.69	60ND2	25C25	67CZ	99.01
66N	25C25	650	33.95	66N	25C25	66C	36.67
66N	25C25	65C	19.02	66N	25C25	60CA	86.34
66N	25C25	61CG	95.07	66N	25C25	61N	72.60
66N	25C25	67N	53.18	een	25 <b>C25</b>	60C	84.37
66N	25C25	60CG	68.93	66N	25025	660	35.32
66N	25C25	60CB	70.27	66N	25C25	61CB	77.27
66 <b>N</b>	25C25	67CG	82.14	6 <b>6</b> N	25C25	67CZ	92.55
650	25C25	66C	65.47	650	25C25	65C	17.60
650	25C25	60CA	65.59	650	25C25	61CG	62.59
650	25C25	61N	40.94	650	25C25	67N	77.29

			TA	BLE XIII	į		
650	25C25	60C	55.85	650	25C25	60CG	66.65
650	25C25	660	68.27	650	25C25	60CB	56.61
650	25C25	61CB	44.60	650	25C25	610D2	71.81
66C	25C25	65C	54.91	66C	25C25	60CA	86.39
66C	25C25	61N	92.94	66C	25C25	67N	17.99
66C	25C25	60C	96.20	66C	25C25	60CG	55.13
66C	25C25	660	13.59	66C	25C25	60CB	68.10
66C	25C25	67CG	47.70	66C	25C25	67CZ	69.56
65C	25C25	60CA	80.60	65C	25C25	61CG	76.13
65C	25C25	61N	58.52	65C	25C25	67N	70.10
65C	25C25	60C	73.03	65C	25C25	60CG	73.62
65C	25C25	660	54.32	65C	25C25	60CB	68.42
65C	25C25	61CB	58.45	65C	25C25	610D2	86.99
60CA	25C25	61CG	66.63	60CA	25C25	61N	34.12
60CA	25C25	590	39.03	60CA	25C25	67N	78.67
60CA	25C25	60C	19.48	60CA	25C25	60CG	33.49
60CA	25C25	660	99.43	60CA	25C25	60CB	18.48
60CA	25C25	61CB	63.35	60CA	25C25	610D2	59.93
61CG	25C25	61N	40.61	61CG	25C25	590	78.90
61CG	25C25	60C	47.83	61CG	25C25	60CG	96.99
61CG	25C25	60CB	78.15	61CG	25C25	61CB	17.99
61CG	25C25	610D2	13.48	61N	25C25	590	66.77
61N	25C25	67 <b>N</b>	95.02	61 <b>N</b>	25C25	60C	16.92
61N	25C25	60CG	57.64	61 <b>N</b>	25C25	60CB	39.04
61N	25C25	61CB	30.85	61 <b>N</b>	25C25	610D2	41.41
590	25C25	67N	92.29	590	25C25	60C	49.98
590	25C25	60CG	56.13	590	25C25	60CB	52.99
590	25C25	61CB	86.68	590	25C25	67CG	96.56
590	25C25	610D2	66.16	67N	25C25	60C	93.00
67N	25C25	60CG	45.25	67N	25C25	660	28.92
67N	25C25	60CB	62.00	67 <b>N</b>	25C25	67CG	38.59
67N	25C25	67CZ	68.17	60C	25C25	60CG	49.75
60C	25C25	60CB	31.34	60C	25C25	61CB	44.00
60C	25C25	610D2	43.11	60 <b>CG</b>	25C25	660	68.70
60CG	25C25	60CB	18.86	60CG	25C25	61CB	88.11
60CG	25C25	67CG	75.30	60CG	25C25	610D2	92.65
660	25C25	60CB	81.00	660	25C25	67CG	47.02
660	25C25	67CZ	61.34	60CB	25C25	61CB	69.79

			TA	BLE XIII			
60CB	25C25	67CG	94.15	60CB	25C25	610D2	74.45
61CB	25C25	610D2	28.84	67CG	25C25	67CZ	33.62
610D1	25C26	650	71.36	610D1	25C26	65C	85.76
610D1	25C26	61CG	11.35	610D1	25C26	61CB	26.51
610D1	25C26	61N	48.33	610D1	25C26	65CA	84.59
610D1	25C26	610D2	15.64	67CE2	25C26	66CA	78.50
67CE2	25C26	6 <b>6N</b>	89.75	67CE2	25C26	67CD2	19.87
67CE2	25C26	66C	60.32	67CE2	25C26	67CZ	10.65
67CE2	25C26	660	56.85	66CA	25C26	66N	21.30
66CA	25C26	650	40.66	66CA	25C26	65C	35.52
66CA	25C26	67CD2	62.19	66CA	25C26	61CG	97.24
66CA	25C26	66C	18.45	66CA	25C26	61CB	83.20
66CA	25C26	61 <b>N</b>	61.53	66CA	25C26	67CZ	84.37
66CA	25C26	65CA	50.10	66CA	25C26	660	28.03
66N	25C26	650	32.82	66N	25C26	65C	19.16
66N	25C26	67CD2	77.48	66N	25C26	61CG	94.77
66N	25C26	66C	32.67	6 <b>6</b> N	25C26	61CB	77.42
66N	25C26	61N	65.23	66N	25C26	67CZ	<b>92.40</b>
66N	25C26	65CA	29.97	6 <b>6N</b>	25C26	660	32.92
650	25C26	65C	17.65	650	25C26	61CG	61.95
650	25C26	66C	58.76	650	25C26	61CB	44.89
650	25C26	61N	36.15	650	25C26	65CA	29.75
650	25C26	610D2	69.17	650	25C26	660	64.19
65C	25C26	67CD2	96.05	65C	25C26	61CG	77.55
65C	25C26	6 <b>6C</b>	50.78	65C	25C26	61CB	59.53
65C	25C26	61N	53.71	65C	25C26	65CA	16.87
65C	25C26	610D2	85.83	65C	25C26	660	52.07
67CD2	25C26	66C	45.30	67CD2	25C26	67CZ	29.89
67CD2	25C26	660	46.56	61CG	25C26	61CB	18.38
61CG	25C26	61N	36.98	61CG	25C26	65CA	79.04
61CG	25C26	610D2	11.28	66C	25C26	61 <b>N</b>	78.34
66C	25C26	67CZ	65.93	66C	25C26	65CA	62.43
66C	25C26	660	14.16	61CB	25C26	61N	29.95
61CB	25C26	65CA	60.78	61CB	25C26	610D2	28.51
61N	25C26	65CA	64.34	61N	25C26	610D2	38.83
51N	25C26	660	89.56	67CZ	25C26	660	59.87
55CA	25C26	610D2	89.18	65CA	25C26	660	59.63
510D1	25C27	61CG	5.16	610D1	25C27	610D2	15.09

			ТА	BLE XIII			•
67CE2	25C27	67CD2	15.74	67CE2	25C27	670H	28.29
61CG	25C27	610D2	12.69	67CD2	25C27	670H	44.01
610D1	25C28	61CG	9.27	610D1	25C28	2640H2	52.17
610D1	25C28	610D2	23.20	67CE2	25C28	67CD2	15.49
61CG	25C28	2640H2	46.24	61CG	25C28	610D2	14.74
2640H2	25C28	610D2	33.06	66N	25C29	65C	24.86
66N	25C29	650	39.60	66N	25C29	66CA	24.12
66N	25C29	65CA	41.74	66N	25C29	67CE2	94.22
66N	25C29	640	81.94	66N	25C29	66C	34.73
66N	25C29	660	37.60	66N	25C29	65N	53.80
66N	25C29	67CD2	79.01	66N	25C29	61CG	98.70
66N	25C29	64C	70.01	66N	25C29	61CB	84.65
65C	25C29	650	20.60	65C	25C29	66CA	42.11
65C	25C29	65CA	24.93	65C	25C29	610D1	91.42
65C	25C29	640	61.11	65C	25C29	66C	58.05
65C	25C29	660	62.45	65C	25C29	65N	31.53
65C	25C29	61CG	79.83	65C	25C29	64C	47.67
65C	25C29	61CB	63.17	650	25C29	66CA	46.14
650	25C29	65CA	39.66	650	25C29	610D1	71.78
650	25C29	640	64.40	650	25C29	66C	65.45
650	25C29	660	74.94	650	25C29	65N	37.56
650	25C29	61CG	60.19	650	25C29	64C	50.20
650	25C29	61CB	45.06	66CA	25C29	65CA	64.24
66CA	25C29	67CE2	75.91	66CA	25C29	66C	19.63
66CA	25C29	660	32.86	66CA	25C29	65N	73.62
66CA	25C29	67CD2	59.00	66CA	25C29	61CG	93.56
66CA	25C29	64C	89.77	66CA	25C29	61CB	85.67
65CA	25C29	610D1	97.52	65CA	25C29	640	40.31
65CA	25C29	66C	76.17	65CA	25C29	660	74.54
65CA	25C29	65N	15.70	65 <b>CA</b>	25C29	61CG	87.20
65CA	25C29	64C	29.78	6 <b>5CA</b>	25C29	61CB	68.64
610D1	25C29	640	78.88	610 <b>D</b> 1	25C29	65N	83.10
610D1	25C29	61CG	11.60	610D1	25C29	64C	76.76
610D1	25C29	61CB	<b>29</b> .29	67CE2	25C29	66C	59.68
67CE2	25C29	660	59.54	67CE2	25C29	67CD2	17.43
640	25C29	65N	29.62	640	25C29	61CG	73.88
640	25C29	64C	14.26	640	25C29	61CB	59.65
66C	25C29	660	16.37	66C	25C29	65N	88.46

			TA	BLE XIII			
66C	25C29	67CD2	44.33	660	25C29	65N	89.10
660	25C29	67CD2	48.06	65N	25C29	61CG	73.50
65N	25C29	64C	16.31	65N	25C29	61CB	55.08
61CG	25C29	64C	69.32	61CG	25C29	61CB	18.57
64C	25C29	61CB	52.55	66N	25030	67CE2	95.44
66N	25030	65C	20.14	66N	25030	660	41.64
66N	25030	66CA	20.76	66N	25030	65CA	36.46
6 <b>6</b> N	25030	6 <b>6</b> C	35.12	66N	25030	67CD2	78.01
66N	25030	650	28.77	66N	25030	640	70.26
67CE2	25030	660	66.02	67CE2	25030	66CA	75.33
67CE2	25030	6 <b>6</b> C	62.32	67CE2	25030	67CD2	17.44
67CE2	25030	67CZ	17.35	67CE2	25030	670H	32.31
65C	25030	660	61.78	65C	25030	66CA	35.78
65C	25030	65CA	21.92	65C	25030	66C	54.08
65C	25030	67CD2	93.11	65C	25030	650	13.87
65C	25030	640	51.57	660	25030	66CA	34.58
660	25030	65CA	74.20	660	25030	66C	16.71
660	25030	67CD2	51.17	660	25030	67CZ	70.40
660	25030	670H	86.34	660	25030	650	68.48
66CA	25030	65CA	55.92	66CA	25030	66C	20.67
66CA	25030	67CD2	58.02	66CA	25030	67CZ	88.15
66CA	25030	650	36.77	66CA	25030	640	87.19
65CA	25030	66C	71.35	65CA	25030	650	31.68
65CA	25030	640	35.13	66C	25030	67CD2	45.28
66C	25030	67CZ	71.68	66C	25030	670H	89.19
66C	25030	650	57.23	67CD2	25030	67CZ	31.46
67CD2	25030	670H	48.52	67CD2	25030	650	89.19
67CZ	25030	670H	17.73	550	25030	640	52.71
6 <b>6N</b>	25C31	65CA	39.43	6 <b>6</b> N	25C31	65C	19.88
66N	25C31	660	41.44	66N	25C31	66CA	16.42
66N	25C31	640	73.76	6 <b>6N</b>	25C31	66C	31.86
66N	25C31	650	24.99	6EN	25C31	67CE2	78.02
6 <b>6</b> N	25C31	65N	43.58	65CA	25C31	65C	22.92
65CA	25C31	660	78.51	65CA	25C31	66CA	54.89
65CA	25C31	640	37.80	65CA	25C31	66C	71.14
65CA	25C31	650	28.31	65CA	25C31	65N	10.41
65C	25C31	660	61.32	65C	25C31	66CA	33.07
65C	25C31	640	54.01	65C	25C31	660	51.04

			TA	ABLE XIII			
65C	25C31	650	9.78	65C	25C31	67CE2	90.50
65C	25C31	65N	30.23	660	25C31		32.38
660	25C31	66C	14.52	660	25C31	650	65.09
660	25C31	67CE2	56.24	660	25C31	65N	88.60
66CA	25C31	640	86.14	66CA	25C31	66C	19.15
66CA	25C31	650	33.97	66CA	25C31	67CE2	61.64
66CA	25C31	65N	63.22	640	25C31	650	52.33
640	25C31	65N	27.39	66C	25C31	650	53.03
66C	25C31	67CE2	52.59	66C	25C31	65N	80.44
650	25C31	67CE2	85.62	650	25C31	65N	32.85
65CA	25032	640	42.64	65CA	25032	66N	34.36
65CA	25032	65C	19.66	65CA	25032	64C	30.83
65CA	25032	65N	14.60	640	25032	66N	73.09
640	25032	65C	55.36	640	25032	64C	12.34
640	25032	65N	28.48	6 <b>6N</b>	25032	65C	17.74
66N	25032	64C	63.03	6 <b>6N</b>	25032	65N	48.15
65C	25032	64C	45.59	65C	25032	65N	31.63
64C	25032	65N	16.37	660	25C33	66N	39.03
660	25C33	65CA	71.47	660	25C33	25SG	98.45
660	25C33	65C	53.71	660	25C33	66C	9.09
66N	25C33	65CA	32.53	6 <b>6</b> N	25C33	25SG	95.78
66N	25C33	65C	14.69	6 <b>6N</b>	25C33	66C	30.21
1610	25C33	161C	15.33	1610	25C33	25SG	60.66
65CA	25C33	25SG	88.64	65CA	25C33	65C	18.08
65CA	25C33	66C	62.73	161C	25C33	25SG	68.01
25 <i>S</i> G	25C33	65C	94.64	65C	25C33	66C	44.83
660	25C34	163CB	85.75	660	25C34	66N	32.40
161C	25C34		16.73	161C	25C34	162N	17.83
161C	25C34		62.24	161C	25C34	162CA	31.39
161C	25C34	162C	47.88	161C	25C34	25 <b>S</b> G	73.05
161C	25C34	161CA	18.94	161C	25C34	163CB	92.63
1610	25C34	162N	30.34	1610	25C34	163N	65.51
1610	25C34	162CA	35.76	1610	25C34	162C	54.67
1610	25C34	25 <b>SG</b>	62.53	1610	25C34	161CA	30.82
1610	25C34	1.63CB	95.27	162X	25C34	163N	46.66
162N	25C34	162CA	18.35	162N	25C34	162C	31.03
162N	25C34	25 <i>S</i> G	69.25	162N	25C34	161CA	31.09
162N	25C34	163CB	76.34	163N	25C34	162CA	31.03

		Т.	ABLE XIII		
163N	25C34 162C	16.65	163N	25C34 25SG	45.82
163N	25C34 161CA	77.75	163N	25C34 163CB	30.49
162CA	25C34 162C	19.01	162CA	25C34 25SG	52.46
162CA	25C34 161CA	48.32	162CA	25C34 163CB	61.52
162C	25C34 25SG	55.74	162C	25C34 161CA	61.81
162C	25C34 163CB	45.32	25 <i>S</i> G	25C34 161CA	91.65
25SG	25C34 163CB	53.96	25SG	25C34 66N	87.39
660	25C35 66C	2.57	660	25C35 163CB	96.66
660	25C35 26CB	46.90	660	25C35 66N	29.09
660	25C35 67CA	31.24	660	25C35 68SD	77.53
660	25C35 67CD1	62.23	660	25C35 67CE1	72.23
66C	25C35 163CB	98.61	66C	25C35 26CB	49.23
66C	25C35 66N	30.32	66C	25C35 67CA	29.98
66C	25C35 68SD	78.11	66C	25C35 67CD1	59.82
66C	25C35 67CE1	69.68	163CB	25C35 26CB	51.66
163CB	25C35 67CA	90.92	163CB	25C35 209CD2	95.04
163CB	25C35 68SD	41.03	26CB	25C35 66N	52.74
26CB	25C35 67CA	56.99	26CB	25C35 68SD	53.93
26CB	25C35 67CD1	97.81	66N	25C35 67CA	60.27
6 <b>6N</b>	25C35 67CD1	84.58	6 <b>6N</b>	25C35 67CE1	89.12
67CA	25C35 209CD2	76.21	67CA	25C35 68SD	56.79
67CA	25C35 67CD1	40.85	67CA	25C35 67CE1	55.83
209CD2	25C35 68SD	66.55	209CD2	25C35 67CD1	48.98
209CD2	25C35 67CE1	49.09	68SD	25C35 67CD1	78.05
68SD	25C35 67CE1	92.43	67CD1	25C35 67CE1	16.13
660	25C36 68SD	96.83	660	25C36 26CB	52.43
660	25C36 66C	6.57	660	25C36 67CA	36.71
660	25C36 26CX	70.05	68SD	25C36 163CB	57.33
68SD	25C36 163CA	70.03	68SD	25C36 134CB	84.74
68SD	25C36 163N	89.11	68SD	25C36 26CB	67.27
68SD	25C36 68CE	22.41	68SD	25C36 209CD2	84.13
68SD	25C36 66C	90.48	68SD	25C36 67CA	67.81
68SD	25C36 162C	99.30	68SD	25C36 26CX	59.77
163CB	25C36 163CA	20.06	163CB	25C36 134CB	81.47
163CB	25C36 163N	34.18	163CB	25C36 26CB	62.38
163CB	25C36 68CE	60.98	163CB	25C36 162C	48.31
163CB	25C36 26CX	44.78	163CA	25C36 134CB	65.51
163CA	25C36 163N	19.48	163CA	25C36 26CB	80.86

		Т	ABLE XIII	Ī	•
163CA	25C36 68CE	65.92		25C36 162C	30.24
163CA	25C36 26CX	63.59	134CB	25C36 163N	69.86
134CB	25C36 68CE	62.66	134CB	25C36 209CD2	53.95
134CB	25C36 162C	59.74	163N	25C36 26CB	84.29
163N	25C36 68CE	85.05	163N	25C36 162C	15.50
163N	25C36 26CX	69.23	26CB	25C36 68CE	88.83
26CB	25C36 66C	50.37	26CB	25C36 67CA	61.54
26CB	25C36 162C	98.98	26CB	25C36 26CX	17.62
68CE	25C36 209CD2	68.72	68CE	25C36 67CA	82.83
68CE	25C36 162C	90.28	68CE	25C36 26CX	78.81
209CD2	25C36 67CA	82.71	66C	25C36 67CA	30.72
66C	25C36 26CX	67.86	67CA	25C36 26CX	75.01
162C	25C36 26CX	84.54	660	25C37 67CE1	76.69
660	25C37 67CZ	72.75	660	25C37 67CD1	64.04
660	25C37 67OH	86.40	660	25C37 67CE2	57.39
660	25C37 67CG	48.64	67CE1	25C37 209CD2	60.17
67CE1	25C37 67CZ	18.76	67 <b>CE</b> 1	25C37 67CD1	18.19
67CE1	25C37 67OH	31.37	67CE1	25C37 67CE2	29.99
67CE1	25C37 67CG	29.19	209CD2	25C37 67CZ	78.42
209CD2	25C37 67CD1	57.12	209CD2	25С37 67ОН	84.22
209CD2	25C37 67CE2	89.26	209CD2	25C37 134CB	51.72
209CD2	25C37 67CG	70.82	67CZ	25C37 67CD1	31.75
67CZ	25С37 67ОН	17.50	67CZ	25C37 67CE2	16.32
67CZ	25C37 67CG	34.21	67CD1	25С37 67ОН	47.81
67CD1	25C37 67CE2	34.53	67CD1	25C37 67CG	15.87
670H	25C37 160O	98.96	670H	25C37 67CE2	29.39
670H	25C37 67CG	51.71	1600	25C37 134CB	78.01
67CE2	25C37 67CG	28.65	65CA	25C38 66N	37.26
65CA	25C38 66O	76.23	65CA	25C38 26CD1	57.25
65CA	25C38 65C	19.78	65CA	25C38 23O	54.21
65CA	25C38 224OH2	55.95	€5CA	25C38 26CB	85.45
65CA	25C38 65N	11.53	65CA	25C38 26CG	68.24
25SG	25C38 26CD1	77.09	25 <b>s</b> G	25C38 230	67.47
25 <i>S</i> G	25C38 1610	68.31	25 <b>SG</b>	25C38 224OH2	89.14
25 <b>S</b> G	25C38 26CB	70.06	25 <b>S</b> G	25C38 26CG	76.35
66N	25C38 66O	39.10	66.N	25C38 26CD1	47.09
66N	25C38 65C	17.53	6611	25C38 23O	72.97
66N	25C38 224OH2	92.76	66 <b>N</b>	25C38 26CB	59.41

			TA	BLE XIII			
66N	25C38	65N	48.17	66N	25C38	26CG	48.12
660	25C38	26CD1	61.41	660	25C38	65C	56.61
660	25C38	26CB	46.70	660	25C38	65N	87.28
660	25C38	26CG	50.49	26CD1	25C38	65C	48.65
26CD1	25C38	230	40.99	26CD1	25C38	2240H2	85.80
26CD1	25C38	26CB	32.92	26CD1	25C38	65N	59.48
26CD1	25C38	26CG	15.33	65C	25C38	230	62.21
65C	25C38	224OH2	75.31	65C	25C38	26CB	70.46
65C	25C38	65N	30.67	65C	25C38	26CG	55.38
230	25C38	2240H2	47.84	230	25C38	26CB	69.70
230	25C38	65N	46.94	230	25C38	26CG	55.26
1610	25C38	224OH2	93.60	2240H2	25C38	65 <b>N</b>	44.65
26CB	25C38	65N	90.52	26CB	25C38	26CG	18.20
65N	25C38	26CG	72.50	66N	25039	26CD1	65.73
66N	25039	65CA	47.24	66N	25039	65C	22.10
6 <b>6N</b>	25039	660	47.99	66N	25039	230	95.16
6 <b>6N</b>	25039	26CG	65.86	6 <b>6N</b>	25039	26CB	78.81
6 <b>6</b> N	25039	26NE1	58.35	66N	25039	66CA	14.64
66N	25039	66C	34.63	66N	25039	65N	57.39
6 <b>6N</b>	25039	650	22.72	6 <b>6N</b>	25039	26CX	95.35
66N	25039	26CD2	57.58	26CD1	25039	65CA	77.50
26CD1	25039	65C	65.28	26CD1	25039	660	80.81
26CD1	25039	230	52.52	26CD1	25039	26CG	20.44
26CD1	25039	26CB	42.80	26CD1	25039	25SG	91.26
26CD1	25039	26NE1	14.54	26CD1	25039	66CA	62.36
26CD1	25039	66C	69.41	26CD1	25039	65N	74.38
26CD1	25039	26N	49.70	26CD1	25039	650	61.20
26CD1	25039	23C	59.40	26CD1	25039	26CX	47.75
26CD1		224OH2	98.62	26CD1	25039	26CD2	17.11
65CA	25039	65C	25.89	65CA	25039	6 <b>6</b> 0	94.13
65CA	25039	230	66.29	65CA	25039	26CG	92.00
65CA	25039	26NE1	63.11	65CA	25039	66CA	61.64
65CA	25039	66C	81.81	55CA	25039	65N	12.36
65CA	25039	650	27.30	65CA	25039	23C	72.79
65CA	25039	2240H2	56.03	65CA	25039	26CD2	82.72
65C	25039	660	70.06	65C	25039	230	78.02
65C	25039	26CG	73.56	65C	25039	26CB	92.36
65C	25039	26NE1	53.27	65C	25039	66CA	36.02

		TA	BLE XIII			
65C	25039 66C	56.60	65C	25039	65 <b>N</b>	35.32
65C	25039 650	4.12	65C	25039		86.65
65C	25039 2240H2	81.45	65C	25039		64.11
660	25039 26CG	65.04	660	25039		58.06
660	25039 26NE1	84.63	660	25039	_	35.35
660	25039 66C	15.25	660	25039	26N	91.09
660	25039 650	70.58	660	25039	26CX	73.00
660	25039 26CD2	63.97	230	25039	26CG	71.55
230	25039 26CB	88.75	230	25039	25 <b>S</b> G	73.57
230	25039 26NE1	46.88	230	25039	65N	54.73
230	25039 26N	69.90	230	25039	650	75.20
230	25039 23C	9.35	230	25039	26CX	82.73
230	25039 2240H2	48.87	230	25039	26CD2	69.59
26CG	25039 26CB	23.44	26CG	25039	25 <b>S</b> G	90.54
26CG	25039 26NE1	33.09	26CG	25039	66CA	57.23
26CG	25039 66C	56.85	26CG	25039	65N	91.62
26CG	25039 26N	42.86	26CG	25039	650	69.92
26CG	25039 23C	77.26	26CG	25039	26CX	33.37
26CG	25039 26CD2	9.50	26CB	25039	25 <b>S</b> G	80.06
26CB	25039 26NE1	56.36	26CB	25039	66CA	66.31
26CB	25039 66C	56.61	26CB	25039	26 <b>N</b>	34.26
26CB	25039 650	89.29	26CB	25039	23C	92.00
26CB	25039 26CX	17.14	26CB	25039	26CD2	31.88
25SG	25039 26N	48.14	25 <b>S</b> G	25039	23C	65.56
25 <i>S</i> G	25039 26CX	62.98	25 <i>S</i> G	25039	2240H2	85.27
25SG	25039 26CD2	98.66	26NE1	25039	66CA	58.87
26NE1	25039 66C	71.02	26NE1	25039	65N	59.89
26NE1	25039 26N	63.22	26 <b>NE1</b>	25039	650	49.16
26NE1	25039 23C	55.32	26NE1	25039	26CX	62.29
26NE1	25039 2240H2	88.14	26NE1	25039	26CD2	26.49
66CA	25039 66C	20.82	66CA	25039	65N	71.23
66CA	25039 26N	98.43	66CA	25039	650	35.79
66CA	25039 26CX	83.32	65CA	25039	26CD2	50.40
66C	25039 65N	91.92	66C	25039	26N	90.87
66C	25039 650	56.59	66C	25039	26CX	73.34
66C	25039 26CD2	53.53	65N	25039	650	35.54
65N	25039 23C	60.74	65N	25039 2	2240H2	46.37
65N	25039 26CD2	82.99	2611	25039	23C	68.72

			T.	ABLE XIII			
26N	25039	26CX	18.10	26N	25039	26CD2	51.68
650	25039	23C	84.01	650	25039	2240H2	81.91
650	25039	26CD2	60.44	23C	25039	26CX	83.51
23C	25039	2240H2	45.98	23C	25039	26CD2	76.48
26CX	25039	26CD2	42.86	25SG	25 <b>N4</b> 0		86.06
25 <b>S</b> G	25 <b>N4</b> 0	230	74.10	25SG	25 <b>N4</b> 0	161C	85.73
25SG	25 <b>N4</b> 0	23C	68.83	25SG	25N40	23CA	76.96
25 <b>S</b> G	25N40	162CA	56.65	25SG	25 <b>N4</b> 0	25CB	6.53
25 <b>S</b> G	25 <b>N4</b> 0	26CD1	72.41	1610	25 <b>N4</b> 0	161C	12.75
1610	25 <b>N4</b> 0	162CA	33.97	1610	25N40	25CB	90.20
2240H2	25 <b>N4</b> 0	230	52.97	2240H2	25 <b>N</b> 40	65CA	57.51
224OH2	25N40	23C	49.02	2240H2	25 <b>N4</b> 0	23CA	35.43
2240H2	25 <b>N4</b> 0	26CD1	85.10	230	25N40	65CA	51.31
230	25 <b>N4</b> 0	23C	13.96	230	25 <b>N4</b> 0	23CA	29.87
230	25 <b>N4</b> 0	25CB	68.50	230	25 <b>N4</b> 0	26CD1	37.36
65CA	25 <b>N4</b> 0	23C	62.89	65CA	25 <b>N4</b> 0	23CA	68.80
65CA	25 <b>N4</b> 0	26CD1	48.33	161C	25 <b>N4</b> 0	162CA	29.53
161C	25N40	25CB	90.88	23C	25 <b>N4</b> 0	23CA	18.22
23C	25N40	25CB	62.59	23C	25 <b>N4</b> 0	26CD1	49.62
23CA	25 <b>N4</b> 0	25CB	70.43	23CA	25N40	26CD1	67.09
162CA	25 <b>N4</b> 0	25CB	62.21	25CB	25 <b>N4</b> 0	26CD1	70.23
25SG	25 <b>N4</b> 1	23C	99.66	25SG	25 <b>N4</b> 1	25CB	18.83
25 <i>\$</i> G	25N41	25N	49.91	25SG	25N41	24N	86.40
25 <i>S</i> G	25 <b>N4</b> 1	1610	75.97	25 <i>S</i> G	25N41	26CD1	82.58
25SG	25N41	25CA	32.30	25 <i>S</i> G	25N41	26N	41.60
230	25N41	2240H2	68.49	230	25N41	23C	20.89
230	25N41	23CA	42.41	230	25 <b>N4</b> 1	25CB	89.72
230	25 <b>N4</b> 1	25N	55.16	230	25N41	24N	28.91
230	25N41	65CA	55.11	230	25N41	26CD1	40.72
230	25N41	25CA	71.69	230	25 <b>N4</b> 1	26N	66.61
230	25N41	23N	44.84	2240H2	25N41	23C	65.79
224OH2	25N41	23CA	48.09	2240H2	25 <b>N4</b> 1	24N	78.14
2240H2	25N41	65CA	60.11	224042	25N41	26CD1	98.71
2240H2	25N41	23N	46.92	23C	25N41	23CA	25.44
23C	25N41	25CB	82.83	23C	25N41	25N	49.85
23C	25N41	24N	13.60	23C	25N41	65CA	73.03
23C	25N41	26CD1	59.09	23C	25 <b>N41</b>	25CA	67.43
23C	25N41	26N	73.57	23C	25 <b>N4</b> 1	23N	27.92

			TA	BLE XIII			
23CA	25N41	25CB	95.07	23CA	25N41	25N	68.21
23CA	25 <b>N4</b> 1	24N	33.15	23CA	25N41	65CA	80.89
23CA	25 <b>N4</b> 1	26CD1	83.00	23CA	25N41	25CA	84.37
23CA	25N41	26N	97.47	23CA	25N41	23N	2.51
25CB	25 <b>N4</b> 1	25N	34.64	25CB	25N41	24N	69.27
25CB	25N41	1610	90.28	25CB	25N41	26CD1	80.97
25CB	25N41	25CA	18.50	25CB	25N41	26N	41.33
25CB	25 <b>N4</b> 1	23N	95.93	25N	25N41	24N	36.95
25N	25N41	26CD1	57.65	25N	25N41	25CA	17.61
25N	25N41	26N	34.79	25N	25N41	23 <b>N</b>	69.91
24N	25N41	65CA	83.83	24N	25N41	26CD1	59.64
24N	25N41	25CA	54.36	24N	25 <b>N4</b> 1	26N	64.38
24N	25N41	23N	35.31	65CA	25N41	26CD1	50.18
65CA	25N41	26N	90.57	65CA	25N41	23N	82.10
26CD1	25N41	25CA	64.68	26CD1	25 <b>N4</b> 1	26N	41.23
26CD1	25 <b>N4</b> 1	23N	85.46	25CA	25N41	26N	29.42
25CA	25N41	23N	85.82	26N	25N41	23N	99.68
660	25 <b>N4</b> 2	66N	49.20	660	25 <b>N4</b> 2	65CA	85.32
660	25N42	66C	14.31	660	25N42	65C	66.36
660	25N42	66CA	34.57	660	25N42	67CE2	59.50
66N	25N42	65CA	37.29	66N	25N42	66C	37.91
66N	25N42	65C	17.17	66N	25N42	66CA	18.36
6 6N	25N42	67CE2	79.25	65CA	25N42	66C	75.14
65CA	25N42	65C	21.61	65CA	25N42	66CA	55.24
66C	25N42	65C	54.69	66C	25N42	66CA	21.22
66C	25N42	67CE2	55.45	65C	25N42	66CA	34.04
65C	25 <b>N4</b> 2	67CE2	88.07	66CA	25N42	67CE2	63.12

## **TABLE XIV**

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
184CB	25C1	184CG	19.63	184CB	25C1	1840	37.95
184CB	25C1	184CD2	33.99	184CB	25C1	184CE3	43.05
184CB	25C1	188CD1	60.59	184CG	25C1	1840	56.47
184CG	25C1	184CD2	18.11	184CG	25C1	184CE3	33.19
184CG	25C1	188CD1	68.63	1840	25C1	184CD2	71.92
1840	25C1	184CE3	78.39	1840	25C1	188CD1	69.15
184CD2	25C1	184CE3	17.41	184CD2	25C1	188CD1	62.78
184CE3	25C1	188CD1	48.96	1840	25C2	184CB	47.37
1840	25C2	184C	14.60	1840	25C2	184CG	67.03
1840	25C2	184CA	35.09	1840	25C2	184CD2	79.86
1840	25C2	184CD1	72.22	1840	25C2	18ND2	51.81
184CB	25C2	184C	37.41	184CB	25C2	184CG	20.57
184CB	25C2	184CA	21.23	184CB	25C2	184CD2	32.66
184CB	25C2	184CD1	31.54	184CB	25C2	18ND2	83.90
184C	25C2	184CG	55.27	184C	25C2	184CA	21.27
184C	25C2	184CD2	69.82	184C	25C2	184CD1	58.47
184C	25C2	18ND2	51.06	184CG	25C2	184CA	35.04
184CG	25C2	184CD2	16.66	184CG	25C2	184CD1	15.49
184CG	25C2	18ND2	93.96	184CA	25C2	184CD2	50.81
184CA	25C2	184CD1	37.24	184CA	25C2	18ND2	62.72
184CD2	25C2	184CD1	26.96	184CD1	25C2	18ND2	86.49
1840	25C3	184CB	45.92	1840	25C3	184CA	37.85
1840	25C3	184C	17.54	1840	25C3	180D1	72.66
1840	25C3	184CG	67.51	1840	25C3	184CD1	78.54
1840	25C3	18ND2	59.83	1840	25C3	18CG	62.25
1840	25C3	184CD2	77.25	1.84CB	25C3	184CA	23.15
184CB	25C3	184C	38.05	184CB	25C3	180D1	86.49
184CB	25C3	184CG	21.78	184CB	25C3	184CD1	36.65
184CB	25C3	18ND2	94.41	184CB	25C3	18CG	86.35
184CB	25C3	184CD2	32.23	184CA	25C3	184C	22.46
184CA	25C3	180D1	63.82	184CA	25C3	184CG	37.27
184CA	25C3	184CD1	42.13	184CA	25C3	18ND2	72.53
184CA	25C3	18CG	63.26	184CA	25C3	184CD2	51.60
184C	25C3	18001	61.36	184C	25C3	184CG	57.47
184C	25C3	184CD1	64.57	184C	25C3	18ND2	57.87

			TA	BLE XIV			
184C	25C3	18CG	54.50	184C	25C3	184CD2	70.12
180D1	25C3	184CG	91.25	180D1	25C3	184CD1	79.96
180D1	25C3	18ND2	31.05	180D1	25C3	18CG	15.09
184CG	25C3	184CD1	18.78	184CG	25C3	18CG	96.39
184CG	25C3	184CD2	15.20	184CD1	25C3	18CG	88.85
184CD1	25C3	184CD2	27.59	18ND2	25C3	18CG	16.95
184CD1	25C4	184CG	19.46	184CD1	25C4	184CB	35.62
184CD1	25C4	180D1	77.64	184CD1	25C4	200	86.16
184CD1	25C4	184CA	39.47	184CD1	25C4	184NE1	16.76
184CD1	25C4	184CD2	28.36	184CD1	25C4	1840	68.14
184CD1	25C4	184CE2	25.97	184CG	25C4	184CB	20.21
184CG	25C4	180D1	84.29	184CG	25C4	184CA	33.31
184CG	25C4	184NE1	29.37	184CG	25C4	184CD2	16.86
184CG	25C4	1840	56.12	184CG	25C4	184CE2	26.96
184CB	25C4	180D1	74.62	184CB	25C4	184CA	19.60
184CB	25C4	184NE1	48.98	184CB	25C4	184CD2	33.22
184CB	25C4	1840	36.02	184CB	25C4	184CE2	46.70
180D1	25C4	200	64.59	180D1	25C4	184CA	55.04
180D1	25C4	184NE1	91.09	180D1	25C4	1840	58.00
200	25C4	184NE1	82.83	200	25C4	184CE2	96.86
184CA	25C4	184NE1	55.98	184CA	25C4	184CD2	49.56
184CA	25C4	1840	30.00	184CA	25C4	184CE2	59.49
184NE1	25C4	184CD2	27.93	184NE1	25C4	1840	83.70
184NE1	25C4	184CE2	15.77	184CD2	25C4	1840	68.31
184CD2	25C4	184CE2	16.44	1840	25C4	184CE2	82.69
184CG	25 <b>C</b> 5	184CD1	18.11	184CG	25C5	184CD2	18.49
184CG	25C5	184NE1	28.96	184CG	25C5	184CE2	29.05
184CG	25C5	184CB	17.96	184CD1	25C5	184CD2	29.17
184CD1	25C5	184NE1	17.37	184CD1	25C5	184CE2	28.25
184CD1	25C5	184CB	32.12	184CD2	25C5	184NE1	28.48
184CD2	25C5	184CE2	17.52	184CD2	25C5	184CB	32.46
184NE1	25C5	184CE2	16.92	184NE1	25C5	184CB	46.34
184CE2	25C5	184CB	46.48	184CD2	25C6	184CG	18.41
184CD2	25C6	184CE3	17.57	J.84CD2	25C6	184CB	32.70
184CD2	25C6	184CE2	16.35	184CD2	25C6	184CD1	26.98
184CG	25C6	184CE3	33.10	184CG	25 <b>C</b> 6	184CB	18.26
184CG	25 <b>C</b> 6	184CE2	27.66	184CG	25C6	184CD1	15.78
184CE3	25C6	184CB	40.99	184CE3	25C6	1430E1	88.61
184CE3	25C6	184CE2	29.17	184CE3	25C6	184CD1	44.38
184CB	25C6	184CE2	45.38	184CB	25C6	184CD1	30.55
184CE2	25C6	184CD1	26.20	200	25C7	20C	19.21
200	25C7	20CA	39.07	200	25 <b>C</b> 7	20N	45.05
200	25C7	180D1	87.05	200	25C7	19CG	55.64
200	25C7	21NE2	74.51	200	25C7	21N	24.68

				BLE XIV			
200	25C7	18CG	94.59	200	25C7	19C	40.56
20C	25C7	20CA	23.55	20C	25C7	20N	38.22
20C	25C7	180D1	77.28	20C	25C7	19CG	67.04
20C	25C7	21NE2	71.32	20C	25C7	21N	11.95
20C	25C7	18CG	83.17	20C	25C7	19C	38.66
20CA	25C7	20 <b>N</b>	21.87	20CA	25 <b>C</b> 7	180D1	54.50
20CA	25C7	19CG	66.31	20CA	25C7	21NE2	87.33
20CA	25C7	21N	31.76	20CA	25C7	18CG	59.80
20CA	25C7	19C	28.41	20N	25C7	180D1	42.02
20N	25C7	184CD1	96.35	20N	25C7	19CG	47.93
20N	25C7	21N	49.41	20N	25C7	18CG	49.82
20N	25C7	19C	10.08	20N	25C7	184CA	86.28
180D1	25 <b>C7</b>	184CD1	83.02	180D1	25C7	19CG	66.42
180D1	25C7	21N	86.24	180D1	25C7	184NE1	98.06
180D1	25C7	184CG	83.82	180D1	25C7	18CG	8.98
180D1	25C7	19C	48.05	180D1	25C7	184CA	54.56
184CD1	25C7	19CG	55.25	184CD1	25C7	184NE1	16.81
184CD1	25C7	184CG	16.56	184CD1	25C7	18CG	85.33
184CD1	25C7	19C	89.88	184CD1	25C7	184CA	36.29
19CG	25C7	21N	77.73	19CG	25C7	184NE1	56.16
19CG	25C7	184CG	70.76	19CG	25C7	18CG	74.86
19CG	25C7	19C	38.66	19CG	25C7	184CA	68.46
21NE2	25C7	21N	59.46	21N	25C7	18CG	91.06
21N	25C7	19C	50.55	184NE1	25C7	184CG	28.00
184NE1	25C7	19C	94.36	184NE1	25C7	184CA	52.97
184CG	25C7	18CG	83.54	184CG	25C7	184CA	30.15
18CG	25C7	19C	56.53	18CG	25C7	184CA	53.47
19C	25C7	184CA	85.50	200	2508	19CG	74.18
200	2508	20C	14.24	200	2508	20N	45.16
200	2508	19CD	84.08	200	2508	20CA	33.67
200	2508	180D1	81.17	200	2508	19C	43.87
200	2508	21N	14.62	200	2508	19CB	65.10
200	2508	19NE2	73.97	1.9CG	2508	184CD1	71.14
19CG	2508	20C	79.40	19CG	2508	184NE1	74.25
19CG	2508	20N	55.59	19CG	2508	19CD	22.00
19CG	2508	20CA	73.82	19CG	2508	190E1	34.44
19CG	2508	184CG	85.17	19CG	2508	180D1	70.42
19CG	2508	184CE2	88.06	1.9CG	2508	19C	42.61
19CG	2508	21N	87.25	1.9CG	2508	19CB	11.43
19CG	2508	19NE2	28.77	19CG	2508	1830	42.56
19 <b>C</b> G	2508	184CD2	93.05	184CD1	2508	184NE1	21.99
184CD1	2508	19CD	64.82	184CD1	2508	1.90E1	49.70
184CD1	2508	184CG	15.65	134CD1	2508	130D1	83.93
184CD1	2508	184CE2	27.56	184CD1	2508	19CB	80.03

			T	ABLE XIV			
184CD1	2508	19NE2	77.46	184CD1	2508	1830	43.29
184CD1	2508	184CD2	22.85	20C	2508	20N	37.96
20C	2508	19CD	93.27	20C	2508	20CA	21.59
20C	2508	180D1	70.08	20C	2508	19C	41.74
20C	2508	21N	10.44	20C	2508	19CB	68.90
20C	2508	19NE2	85.00	184NE1	2508	19CD	59.98
184NE1	2508	190E1	43.21	184NE1	2508	184CG	31.35
184NE1	2508	184CE2	13.82	184NE1	2508	19CB	85.19
184NE1	2508	19NE2	69.30	184NE1	2508	1830	60.22
184NE1	2508	184CD2	24.73	20N	2508	19CD	76.55
20N	2508	20CA	20.72	20N	2508	190E1	89.93
20N	2508	180D1	38.33	20N	2508	19C	14.64
20N	2508	21N	48.38	20N	2508	19CB	44.35
20N	2508	19NE2	76.55	20N	2508	1830	65.49
19CD	2508	20CA	92.98	19CD	2508	190E1	16.77
19CD	2508	184CG	80.43	19CD	2508	180D1	91.12
19CD	2508	184CE2	73.34	19CD	2508	19C	62.65
19CD	2508	21N	98.59	19CD	2508	19CB	32.44
19CD	2508	19NE2	14.03	19CD	2508	1830	53.18
19CD	2508	184CD2	82.97	20CA	2508	180D1	48.49
20CA	2508	19C	31.25	20CA	2508	21N	31.00
20CA	2508	19CB	62.39	20CA	2508	19NE2	89.43
20CA	2508	1830	85.40	190E1	2508	184CG	65.32
190E1	2508	180D1	95.73	190E1	2508	184CE2	56.63
190E1	2508	19C	77.00	190E1	2508	19CB	45.84
190E1	2508	19NE2	27.83	190E1	2508	1830	50.05
190E1	2508	184CD2	66.42	184CG	2508	180D1	81.85
184CG	2508	184CE2	29.41	184CG	2508	19CB	92.98
184CG	2508	19NE2	93.11	184CG	2508	1830	51.14
184CG	2508	184CD2	15.78	180D1	2508	19C	48.36
180D1	2508	21N	79.10	180D1	2508	19CB	63.32
180D1	2508	19NE2	98.90	180D1	2508	1830	49.03
180D1	2508	184CD2	97.37	184CE2	2508	19CB	98.94
184CE2	2508	19NE2	81.67	184CE2	2508	1830	70.16
184CE2	2508	184CD2	16.41	19C	2508	21N	51.75
19C	2508	19CB	31.18	19C	2508	19NE2	61.93
19C	2508	1830	62.03	21N	2508	19CB	77.34
21N	2508	19NE2	88.58	19CB	2508	19NE2	35.99
19CB	2508	1830	45.66	19NE2	2508	1830	66.29
19NE2	2508	184CD2	93.54	1930	2508	184CD2	64.88
200	25C9	19CG	60.67	200	25C9	19CD	76.38
200	25C9	190E1	92.50	200	25C9	20C	7.58
200	25 <b>C</b> 9	19NE2	72.45	200	25C9	21NE2	67.37
184NE1	25C9	184CD1	20.97	134NE1	25 <b>C</b> 9	19CG	68.64

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			TA	BLE XIV			
184NE1	25C9	19CD	59.83	184NE1	25C9	190E1	43.65
184NE1	25C9	184CE2	16.49	184NE1	25C9	19NE2	71.84
184NE1	25C9	184CG	26.81	184NE1	25C9	184CZ2	28.68
184CD1	25C9	19CG	60.92	184CD1	25C9	19CD	60.57
184CD1	25C9	190E1	48.05	184CD1	25C9	184CE2	31.05
184CD1	25C9	19NE2	75.62	184CD1	25C9	184CG	13.88
184CD1	25C9	184CZ2	46.32	19CG	25C9	19CD	21.52
19CG	25C9	190E1	33.58	19CG	25C9	20C	65.19
19CG	25C9	184CE2	85.10	19CG	25C9	19NE2	31.33
19CG	25C9	184CG	73.53	19CG	25C9	184CZ2	95.37
19CD	25C9	190E1	16.79	19CD	25C9	20C	82.31
19CD	25C9	184CE2	75.59	19CD	25C9	19NE2	15.81
19CD	25C9	184CG	74.44	19CD	25C9	184CZ2	81.71
190E1	25C9	20C	97.96	190E1	25C9	184CE2	59.01
190E1	25C9	19NE2	28.56	190E1	25C9	184CG	61.65
190E1	25C9	184CZ2	65.02	20C	25C9	19NE2	79.44
20C	25C9	21NE2	62.11	184CE2	25C9	19NE2	86.32
184CE2	25C9	184CG	28.77	184CE2	25C9	184CZ2	15.52
19NE2	25C9	184CG	89.47	19NE2	25C9	184CZ2	89.08
184CG	25C9	184CZ2	43.90	184NE1	25010	184CE2	19.56
184NE1	25010	184CD1	19.15	184NE1	25010	200	98.89
184NE1	25010	184CZ2	34.13	184NE1	25010	184CD2	27.67
184NE1	25010	184CG	27.32	184CE2	25010	184CD1	31.59
184CE2	25010	184CZ2	18.04	184CE2	25010	184CD2	16.51
184CE2	25010	184CG	28.75	184CD1	25010	200	88.35
184CD1	25010	184CZ2	49.32	184CD1	25010	184CD2	28.37
184CD1	25010	184CG	15.83	200	25010	184CG	99.14
200	25010	21NE2	63.32	184CZ2	25010	184CD2	30.98
184CZ2	25010	184CG	46.20	184CD2	25010	184CG	17.35
19NE2	25C11	220	38.20	19NE2	25C11	19CD	17.94
19NE2	25C11	200	68.83	19 <b>NE</b> 2	25C11	190E1	29.76
19NE2	25C11	184NE1	67.70	19NE2	25C11	22C	51.21
19NE2	25C11	23CA	50.68	19NE2		19CG	29.78
19NE2	25C11	22N	68.31	19NE2	25C11	23N	56.54
220	25C11	19CD	51.52	220	25C11	200	54.51
220	25C11	190E1	66.66	220	25C11	22C	14.40
220	25C11	23CA	34.19	220	25C11	19CG	50.00
220	25C11	22N	32.97	220	25C11	23N	26.65
19CD	25C11	200	64.43	19CD	25C11	190E1	15.96
19CD	25C11		51.25	1.9CD	25C11	22C	65.67
1900	25C11	23CA	68.61	19CD	25C11	19CG	17.66
19CD	25C11	22N	76.34	19CD	25C11	23N	73.36
200	25C11	190El	76.37	200	25C11		85.38
200	25C11	22C	60.00	200	25C11	23CA	87.97

			TAI	BLE XIV			
200	25C11	19CG	47.41	200	25C11	22N	38.68
200	25C11	23N	75.19	190E1	25C11	184NE1	38.17
190E1	25C11	22C	80.44	190E1	25C11	23CA	78.31
190E1	25C11	19CG	29.35	190E1	25C11	22N	92.26
190E1	25C11	23N	86.19	184NE1	25C11	19CG	53.83
22C	25C11	23CA	29.04	22C	25C11	19CG	64.17
22C	25C11	22N	28.19	22C	25C11	23N	15.42
23CA	25C11	19CG	76.14	23CA	25C11	22N	56.58
23CA	25C11	23N	16.87	19CG	25C11	22N	66.36
19CG	25C11	23N	75.77	22N	25C11	23N	40.54
220	25C12	22C	17.89	220	25C12	22N	39.45
220	25C12	23N	32.43	220	25C12	23CA	38.69
220	25C12	200	56.27	220	25C12	21C	55.84
220	25C12	22CA	31.37	220	25C12	21CA	67.07
220	25C12	19NE2	35.21	22C	25C12	22N	34.05
22C	25C12	23N	18.76	22C	25C12	23CA	33.51
22C	25C12	200	65.66	22C	25C12	21C	49.03
22C	25C12	22CA	18.77	22C	25C12	21CA	64.80
22C	25C12	19NE2	51.68	22N	25C12	23N	49.61
22N	25C12	23CA	67.28	22N	25C12	200	43.43
22N	25C12	21C	16.39	22N	25C12	22CA	18.17
22N	25C12	210E1	71.11	22N	25C12	21CA	30.79
22N	25C12	19NE2	70.87	23N	25C12	23CA	19.64
23N	25C12	200	84.42	23N	25C12	21C	62.12
23N	25C12	22CA	31.69	23N	25C12	21CA	79.89
23N	25C12	19NE2	58.72	23CA	25C12	200	94.95
23CA	25C12	21C	81.09	23CA	25C12	22CA	50.18
23CA	25C12	21CA	98.06	23CA	25C12		50.86
200	25C12	21C	45.48	200	25C12		58.80
200	25C12	210E1	66.04	200	25C12		35.16
200	25C12	19NE2	63.11	21C	25C12		30.96
21C	25C12	210E1	55.27	21C	25C12		18.83
21C	25C12	19NE2	86.29	22CA	25C12		85.66
22CA	25C12	21CA	48.22	22CA			66.48
210E1	25C12	21CA	42.41	21CA	25C12		90.68
210E1	25C13	21CD	12.45	210E1			47.08
210E1	25C13	22N	76.49	210E1			61.27
210E1	25C13		69.05	210E1	25C13		23.98
21CD	25C13		41.21	21CD			71.61
21CD	25C13	21C	58.15	21CD	_		57.98
21CD	25C13		14.96	21CA			30.42
21CA	25C13		18.99	21CA			33.62
21CA	25C13		51.09	22N	25C13		16.45
22N	25C13	200	38.35	22N	25C13	3 21NE2	80.82

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			TA	BLE XIV			
21C	25C13	200	42.37	21C	25C13	21NE2	69.53
200	25C13	21NE2	58.27	210E1	25C14	21C	80.26
210E1	25C14	21CA	58.86	210E1	25C14	22N	95.27
210E1	25C14	21CD	11.97	210E1	25C14	210	83.12
210E1	25C14	21CB	43.94	210E1	25C14	21CG	26.08
210E1	25C14	21NE2	17.70	210E1	25C14	200	71.89
21C	25C14	21CA	23.60	21C	25C14	22N	19.79
21C	25C14	21CD	70.89	21C	25C14	210	18.03
21C	25C14	21CB	37.30	21C	25C14	22CA	32.39
21C	25C14	21CG	54.18	21C	25C14	22C	48.65
21C	25C14	21NE2	78.83	21C	25C14	23N	61.74
21C	25C14	200	44.43	21CA	25C14	22N	36.47
21CA	25C14	21CD	48.38	21CA	25C14	210	36.17
21CA	25C14	21CB	21.86	21CA	25C14	22CA	53.26
21CA	25C14	21CG	33.45	21CA	25C14	22C	65.81
21CA	25C14	21NE2	55.36	21CA	25C14	23N	80.87
21CA	25C14	200	33.68	22N	25C14	21CD	84.38
22N	25C14	210	33.19	22N	25C14	21CB	55.22
22N	25C14	22CA	17.77	22N	25C14	21CG	69.86
22N	25C14	22C	29.90	22N	25C14	21NE2	89.31
22N	25C14	23N	44.46	22N	25C14	200	39.38
21CD	25C14	210	76.50	21CD	25C14	21CB	36.73
21CD	25C14	21CG	18.01	21CD	25C14	21NE2	12.68
21CD	25C14	200	60.03	210	25C14	21CB	39.79
210	25C14	22CA	37.66	210	25C14	21CG	58.51
210	25C14	22C	56.26	210	25C14	21NE2	86.99
210	25C14	23N	65.51	210	25C14	200	62.32
21CB	25C14	22CA	69.70	21CB	25C14	21CG	18.72
21CB	25C14	22C	85.11	21CB	25C14	21NE2	47.76
21CB	25C14	23N	98.97	21CB	25C14	200	53.00
22CA	25C14	21CG	86.05	22CA	25C14	22C	18.65
22CA	25C14	23N	29.45	22CA	25C14	200	54.92
21CG	25C14	22C	99.21	21CG	25C14	21NE2	29.49
21CG	25C14		54.91	22C	25C14	23N	15.74
22C	25C14	200	55.72	21NE2	25C14	200	58.56
23N	25C14	200	70.69	210E1	25C15	21CD	18.02
210E1	25C15	21NE2	36.11	210E1	25C15	21CA	45.66
210E1	25C15	200	74.47	210E1	25C15	21CG	20.94
21CD	25C15	21NE2	20.43	21CD	25C15	21CA	42.29
21CD	25C15	200	63.32	21CD	25C15	21CG	11.31
21NE2	25C15	21CA	56.77	21NE2	25C15	200	66.57
21NE2	25C15	21CG	29.05	21CA	25C15	200	33.79
21CA	25C15	21CG	31.00	200	25C15	21CG	54.47
19NE2	25C16	190E1	34.42	19NE2	25C16	19CD	19.70

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				ABLE XIV			
19NE2	25C16	- <del>-</del>	57.42	19NE2	25C16	184NE1	74.14
19NE2	25C16		38.47	19NE2	25C16	162ND1	77.08
19NE2	25C16		27.65	19NE2	25C16	23N	58.92
19NE2	25C16	_	50.59	190E1	25C16	19CD	18.10
190E1	25C16	23CA	91.00	190E1	25C16	184NE1	41.37
190E1	25C16	220	70.27	190E1	25C16	162ND1	54.22
190E1	25C16	19CG	28.11	190E1	25C16	23N	93.23
190E1	25C16	22C	83.37	190E1	25C16	184CZ2	67.06
19CD	25C16	23CA	76.87	19CD	25C16	184NE1	54.73
19CD	25C16	220	52.37	19CD	25C16	162ND1	70.37
19CD	25C16	19CG	14.40	19CD	25C16	23N	76.48
19CD	25C16	22C	65.58	19CD	25C16	184CZ2	83.17
23CA	25C16	220	36.36	23CA	25C16	19CG	79.64
23CA	25C16	23N	16.38	23CA	25C16	22C	28.95
184NE1	25C16		56.55	184NE1	25C16	19CG	54.16
184NE1	25C16		30.24	220	25C16	19CG	48.64
220	25C16		26.88	220	25C16	22C	13.31
162ND1	25C16		82.32	162ND1	25C16	184CZ2	51.21
19CG	25C16		75.10	19CG	25C16	22C	61.77
19CG	25C16		84.32	23N	25C16	22C	15.32
162ND1	25S17		72.20	162ND1	25S17	184NE1	73.37
162ND1	25517		19.38	162ND1	25 <b>S</b> 17	190E1	64.03
162ND1	25 <b>S</b> 17		73.12	162ND1	25S17	162CG	16.82
162ND1	25517	19CD	76.25	162ND1	25517	19NE2	80.24
162ND1	25817		31.46	162ND1	<b>25</b> S17	184CH2	75.04
162ND1	25517		24.22	162ND1	25517	184CD1	76.19
184CZ2	25S17		40.35	184CZ2	25S17	162CE1	61.98
184CZ2	25S17	190E1	82.50	184CZ2	25 <b>S</b> 17	184CE2	20.39
184CZ2	25517	162CG	64.57	184CZ2	25\$17	19CD	94.35
184CZ2	25517		73.92	184CZ2	25517	184CH2	9.57
184CZ2	25S17		50.06	184CZ2	25 <b>S</b> 17	184CD1	46.22
184NE1	25\$17	162CE1	54.53	184NE1	25 <b>S17</b>	190E1	45.35
184NE1	25517	184CE2	20.06	184NE1		162CG	77.66
	25S17					19NE2	71.53
184NE1		162CB	94.60	184NE1		184CH2	49.77
184NE1		162NE2	51.06	184NE1	25 <b>s</b> 17		6.15
1.62CE1	25517	190E1	48.96	162CE1	25517		57.43
162CE1		162CG	30.44	162 <b>CE1</b>		19CD	62.94
162CE1		19NE2	71.39	162CE1		162CB	48.22
162CE1		184CH2	67.56	162CE1	25\$17		13.10
162CE1	25S17		56.98	190E1		184CE2	64.06
1.90E1		162CG	78.89	190E1		19CD	14.86
190E1		19NE2	29.15			162CB	95.48
190El	<b>25</b> \$17	184CH2	91.95	1.90E1	25 <i>\$</i> 17	162NE2	57.76

			TA	BLE XIV			
190E1	25\$17	184CD1	41.33	184CE2	25\$17	162CG	71.43
184CE2	25\$17	19CD	74.69	184CE2	25S17	19NE2	91.37
184CE2	25S17	162CB	85.35	184CE2	25817	184CH2	29.73
184CE2	25\$17	162NE2	48.91	184CE2	25 <b>S</b> 17	184CD1	25.84
162CG	25S17	19CD	92.05	162CG	25517	19NE2	97.02
162CG	25517	162CB	18.20	162CG	25517	184CH2	64.89
162CG	<b>25</b> S17	162NE2	27.13	162CG	25S17	184CD1	81.91
19CD	25 <b>S</b> 17	19NE2	16.70	19CD	25S17	162NE2	72.44
19CD	25\$17	184CD1	49.68	19NE2	25 <b>S</b> 17	162NE2	82.84
19NE2	25\$17	184CD1	66.32	162CB	25 <b>S</b> 17	184CH2	71.34
162CB	25\$1.7	162NE2	45.10	162CB	25 <b>S</b> 17	184CD1	99.30
184CH2	25\$17	162NE2	54.91	184CH2	25517	184CD1	55.56
162NE2	25S17	184CD1	54.94	19NE2	25N18	23CA	73.04
19NE2	25N18	19CD	18.20	19NE2	25N18	220	43.17
19NE2	25N18		33.64	19NE2	25N18	23C	62.17
19NE2	25N18	25 <i>S</i> G	78.34	19NE2	25N18	23N	70.88
19NE2	25N18	22C	57.87	19NE2	25 <b>N</b> 18	25CB	58.17
19NE2	25N18	24N	45.44	19NE2	25N18	162ND1	81.06
19NE2	25N18		41.89	19NE2	25N18	230	71.35
23CA	25N18		90.63	23CA	25 <b>N</b> 18	220	43.03
23CA	25N18	23C	19.70	23CA	25N18	25 <i>S</i> G	84.26
23CA	25N18	23N	17.82	23CA	25N18	22C	32.08
23CA	25N18	25CB	90.92	23CA	25N18	24N	31.17
23CA	25N18	25N	62.49	23CA	25N18	230	26.79
19CD	25N18	220	55.64	19CD	25N18	190E1	17.64
19CD	25N18	23C	80.36	19CD	25N18	25SG	84.40
19CD	25N18	23N	86.18	19CD	25N18	22C	71.10
19CD	25N18	25CB	60.97	19CD	25N18	24N	63.63
19CD	25N18	162ND1	70.54	19CD	25N18	25N	54.63
19CD	25N18	230	89.15	220	25N18	190E1	73.27
220	25N18	23C	46.87	220	25N18	23N	31.78
220	25N18	22C	15.47	220	25N18	25CB	93.37
220	25N18	24N	37.57	220	25N18	25N	64.91
220	25N18	230	60.03	190E1	25N18	23C	93.60
190E1	25N18	25SG	78.94	190E1	25N18	22C	88.73
190E1	25N18	25CB	54.74	190E1	25N18	24N	77.36
190E1	25N18	162ND1	53.80	1.90E1	25N18	25N	59.44
190E1	25N18	230	99.98	23C	25N18	25SG	66.71
23C	25N18	23N	34.39	23C	25N18	22C	42.39
23C	25N18	25CB	71.24	23C	25N18	24N	16.74
23C	25N18	25N	43.13	23C	25N18	230	13.17
25 <i>S</i> G	25N18	25CB	24.23	25SG	25N18	24N	67.84
25SG	25N18	162ND1	54.04	25 <i>\$</i> G	25N18	25N	40.11
25 <i>S</i> G	25N18	230	57.50	23N	25N18	22C	17.29

			TA	BLE XIV			
23N	25N18	24N	39.13	23N	25N18	25N	73.32
23 <b>N</b>	25N18	230	44.07	22C	25N18	24N	39.37
22C	25N18	25N	71.95	22C	25N18	230	54.65
25CB	25N18	24N	64.90	25CB	25N18	162ND1	43.82
25CB	25N18	25N	30.73	25CB	25N18	230	66.97
24N	25N18	25N	34.26	24N	25N18	230	27.47
162ND1	25N18	25N	73.32	25N	25N18	230	43.35
25SG	25C19	25CB	30.95	25SG	25C19	19NE2	87.88
25SG	25C19	162ND1	72.09	25SG	25C19	23CA	92.18
25SG	25C19	190E1	90.53	25SG	25C19	1610	75.25
25SG	25C19	19CD	92.51	25SG	25C19	25N	44.77
25 <b>S</b> G	25C19	23C	73.06	25 <b>S</b> G	25C19	162CE1	76.49
25SG	25C19	25CA	36.53	25 <b>S</b> G	25C19	230	62.97
25SG	25C19	162CG	75.04	25 <b>S</b> G	25C19	24N	71.77
25 <i>S</i> G	25C19	162CA	61.03	25CB	25C19	19NE2	64.00
25CB	25C19	162ND1	55.38	25CB	25C19	23CA	96.18
25CB	25C19	190E1	5 <b>9</b> .77	25CB	25C19	1610	94.40
25CB	25C19	19CD	63.98	25CB	25C19	25N	33.81
25CB	25C19	23C	76.97	25CB	25C19	162CE1	52.78
25CB	25C19	25CA	15.71	25CB	25C19	230	73.90
25CB	25C19	162CG	63.84	25CB	25C19	24N	67.67
25CB	25C19	162CA	66.83	19NE2	25C19	162ND1	88.64
19NE2	25C19	23CA	60.82	19NE2	25C19	190E1	31.76
19NE2	25C19	19CD	16.34	19NE2	25C19	25N	44.95
19NE2	25C19	23C	56.12	19NE2	25C19	162CE1	73.67
19NE2	25C19	25CA	52.15	19NE2	25C19	230	68.16
19NE2	25C19	24N	41.52	162ND1	25C19	190E1	59.55
162ND1	25C19	1610	71.43	162ND1	25C19	19CD	75.22
162ND1	25C19	25N	86.60	162ND1	25C19	162CE1	15.59
162ND1	25C19	25CA	68.74	162ND1	25C19	162CG	12.11
162ND1	25C19	162CA	39.22	23CA	25C19	190E1	92.31
23CA	25C19	19CD	76.28	23CA	25C19	25N	62.61
23CA	25C19	23C	19.89	23CA	25C19	25CA	81.01
23CA	25C19	230	29.74	23CA	25C19	24N	29.49
190E1	25C19	19CD	16.09	190E1	25C19	25 <b>N</b>	61.21
190E1	25C19	23C	87.07	190E1	25C19	162CE1	43.98
190E1	25C19	25CA	56.66	190E1	25C19	230	97.48
190E1	25C19	162CG	70.72	190E1	25C19	24N	71.88
190E1	25C19		98.02	1610	25C19	162CE1	86.98
1610	25C19	162CG	60.30	1610	25C19	162CA	34.11
19CD	25C19	25N	54.73	19CD	25C19	23C	72.44
19CD	25C19	162CE1	59.74	19CD	25C19	25CA	56.11
19CD	25C19	230	84.17	19CD	25C19	162CG	86.63
19CD	25C19	24N	57.70	25N	25C19	23C	44.11

			TAI	BLE XIV			
25N	25C19	162CE1	79.19	25N	25C19	25CA	18.40
25N	25C19	230	44.87	25N	25C19	162CG	96.59
25N	25C19	24N	33.86	25N	25C19	162CA	99.25
23C	25C19	25CA	62.33	23C	25C19	230	14.57
23C	25C19	24N	15.37	162CE1	25C19	25CA	63.05
162CE1	25C19	162CG	26.96	162CE1	25C19	162CA	54.63
25CA	25C19	230	61.33	25CA	25C19	162CG	78.31
25CA	25C19	24N	52.11	25CA	25C19	162CA	82.37
230	25C19	24N	26.74	162CG	25C19	162CA	30.21
200	25N20	19CD	87.74	200	25N20	19NE2	89.20
200	25N20	19CG	65.65	200	25 <b>N2</b> 0	220	66.51
200	25N20	20C	2.25	200	25N20	184CD1	99.19
200	25N20	22N	43.43	200	25N20	22C	67.94
200	25N20	21CA	31.81	200	25N20	19CB	59.50
19CD	25N20	19NE2	20.94	19CD	25N20	19CG	24.17
19CD	25N20	190E1	18.66	19CD	25N20	220	59.53
19CD	25N20	184NE1	61.57	19CD	25N20	20C	89.98
19CD	25N20	184CD1	58.87	19CD	25N20	22N	90.16
19CD	25N20	22C	72.14	19CD	25 <b>N</b> 20	184CE2	73.87
19CD	25N20	19CB	28.25	19NE2	25 <b>N</b> 20	19CG	38.17
19NE2	25N20	190E1	34.54	19NE2	25 <b>N2</b> 0	220	42.18
19NE2	25N20	184NE1	79.08	19NE2	25N20	20C	91.23
19NE2	25N20	184CD1	79.14	19NE2	25N20	22N	76.95
19NE2	25N20	22C	53.71	19NE2	25N20	184CE2	90.08
19NE2	25N20	19CB	35.63	19CG	25N20	190E1	37.15
19CG	25N20	220	61.25	19CG	25N20	184NE1	67.75
19CG	25N20	20C	67.89	19CG	25N20	184CD1	57.44
19CG	25N20	22N	80.74	19CG	25 <b>N2</b> 0	22C	74.00
19CG	25N20	21CA	93.39	19CG	25N20	184CE2	80.81
19CG	25 <b>N2</b> 0	19CB	11.02	190E1	25 <b>N</b> 20	220	76.29
190E1	25N20	184NE1	44.61	190E1	25 <b>N</b> 20	184CD1	46.08
190E1	25N20	22C	88.22	190E1	25N20	184CE2	56.16
190E1	25N20	19CB	44.63	220	25N20		67.58
220	25N20	22N	36.33	220	25N20		13.03
220	25N20	21CA	65.40	220	25N20		51.11
184NE1	25N20	184CD1	17.40	184NE1	25N20	184CE2	13.10
184NE1	25N20	19CB	78.62	20C	25N20		43.20
20C	25N20	22C	68.43	20C	25N20		30.05
20C	25N20	19CB	61.74	184CD1	25N20		27.05
184CD1	25N20		68.40	22N	25N20		29.81
22N	25N20		29.53	22N	25N20		69.93
22C	25N20	21CA	59.18	22C	25N20		63.62
21CA	25N20	19CB	84.67	184CE2	25N20		91.71
162ND1	25C21		76.59	162ND1	25C21	. 162CE1	18.63

			TA	BLE XIV			
162ND1	25C21	162CG	18.63	162ND1	25C21	1610	88.59
162ND1	25C21	25CB	58.66	162ND1	25C21	162CB	39.41
162ND1	25C21	190E1	66.84	162ND1	25C21	162CA	48.39
162ND1	25C21	19NE2	91.07	162ND1	25C21	19CD	80.09
162ND1	25C21	162NE2	16.42	162ND1	25C21	161C	78.48
162ND1	25C21	184CZ2	60.80	162ND1	25C21	162CD2	15.86
162ND1	25C21	184NE1	63.09	162ND1	25C21	162N	62.96
25 <i>S</i> G	25C21	162CE1	82.30	25SG	25C21	162CG	85.48
25 <b>S</b> G	25C21	1610	75.19	25SG	25C21	25CB	27.93
25SG	25C21	162CB	86.01	25SG	25C21	190E1	84.07
25 <i>S</i> G	25C21	162CA	67.62	25 <i>S</i> G	25C21	19NE2	71.84
25 <i>S</i> G	25C21	19CD	81.26	25 <i>S</i> G	25C21	162NE2	89.59
25 <i>S</i> G	25C21	161C	76.87	25 <i>S</i> G	25C21	162CD2	91.13
25SG	25C21	162N	73.63	162CE1	25C21	162CG	34.59
162CE1	25C21	25CB	58.15	162CE1	25C21	162CB	56.62
162CE1	25C21	190E1	49.16	162CE1	25C21	162CA	67.01
162CE1	25C21	19NE2	75.88	162CE1	25C21	19CD	63.04
162CE1	25C21	162NE2	10.76	162CE1	25C21	161C	97.04
162CE1	25C21		52.78	162CE1	25C21	162CD2	25.92
162CE1	25C21	184NE1	46.57	162CE1	25C21	162N	81.54
162CG	25C21	1610	74.86	162CG	25C21	25CB	73.56
162CG	25C21	162CB	22.12	162CG	25C21	190E1	83.74
162CG	25C21	162CA	37.08	162CG	25C21	19CD	97.60
162CG	25C21	162NE2	27.00	162CG	25C21	161C	64.13
162CG	25C21	184CZ2	58.58	162CG	25C21	162CD2	11.67
162CG	25C21	184NE1	71.27	162CG	25C21	162N	49.24
1610	25C21	25CB	96.45	1610	25C21	162CB	53.58
1610	25C21	162CA	40.31	1610	25C21	161C	10.85
1610	25C21	162CD2	86.16	1610	25C21	162N	25.81
25CB	25C21	162CB	84.13	25CB	25C21	190E1	58.32
25CB	25C21	162CA	73.26	25 <b>CB</b>	25C21	19NE2	55.61
25CB	25C21	19CD	59.29	25CB	25C21	162NE2	67.37
25CB	25C21	161C	94.34	25CB	25C21	162CD2	74.51
25CB		184NE1	89.62	25CB	25C21		85.19
162CB		162CA	21.46	162CB		162NE2	48.98
162CB	25C21		42.74	162CB		184CZ2	70.53
162CB		162CD2	32.77	162CB		184NE1	90.36
162CB	2.5C21		28.88	190E1		19NE2	30.01
190E1		19CD	14.42	190E1		162NE2	57.58
190E1		184CZ2	67.32	190E1		162CD2	74.15
190E1		184NE1	37.69	162CA		162NE2	62.57
162CA	25C21		30.75	162CA		184CZ2	91.75
162CA		162CD2	48.73	162CA	25C21		15.49
19NE2	25C21	19CD	16.41	19NE2	25C21	162NE2	85.63

			7	ABLE XIV	,		
19N	E2 25C2	1 184CZ2	94.21	. 19NE2		184NE1	63.05
190	D 25C2	1 162NE2	71.85		25C21		
19C	D 25C2	1 162CD2	88.39		25C21	184NE1	78.23
162N	E2 25C2:	1 161C	91.07			184CZ2	47.85
162N	E2 25C2	1 162CD2	16.57		_	184NE1	46.06
162N	E2 25C2:	1 162N	75.95		25C21	162CD2	47.02
161C	25C2:	1 162N	15.54			162CD2	75.36
184C2	22 25C2	1 184NE1	30.38			162N	49.06
162CI	25C21		59.61				97.18
25 <i>S</i> G	25C22	25CB	34.32	25SG	25C22	25N	60.72
25 <i>S</i> G	25C22	230	96.62	25SG	25C22	25CA	58.23
25 <i>S</i> G	25C22	1610	88.83	25SG	25C22	19NE2	41.06
25 <i>S</i> G	25C22	24N	94.80	25SG		162ND1	94.39
25 <i>S</i> G	25C22	26N	28.02	25SG	25C22	24C	67.14 63.81
25 <i>S</i> G	25C22	25C	28.81	25SG	25C22	162CA	61.80
25 <i>S</i> G	25C22	26CD1	59.67	25CB	25C22	25N	41.88
25CB	25C22	23C	94.72	25CB	25C22	230	97.61
25CB	25C22	25CA	20.48	25CB	25C22	19NE2	62.01
25CB	25C22	24N	79.22	25CB		162ND1	50.87
25CB	25C22	26N	40.24	25CB	25C22	24C	51.06
25CB	25C22	25C	25.68	25CB		162CA	70.19
25CB	25C22	26CD1	77.04	25N	25C22	23CA	70.13
25N	25C22	23C	53.07	25N	25C22	230	56.89
25N	25C22	25CA	21.75	25N	25C22	19NE2	46.70
25N	25C22	24N	38.64	25N		162ND1	88.45
25N	25C22	26N	36.83	25N	25C22	24C	9.50
25N	25C22	25C	29.43	25N	25C22	26CD1	53.68
23CA	25C22	23C	23.35	23CA	25C22	230	37.19
23CA	25C22	25CA	93.17	23CA	25C22	19NE2	58.05
23CA	25C22	24N	33.59	23CA	25C22	26N	99.96
23CA	25C22	24C	64.82	23CA	25C22	26CD1	78.54
23C	25C22	230	18.81	23C	25C22	25CA	74.82
23C	25C22	19NE2	57.79	23C	25C22	24N	16.66
23C	25C22	26N	76.83	23C	25C22	24C	44.46
23C	25C22	25C	78.94	23C	25C22	26CD1	57.31
230	25C22	25CA	77.22	230	25C22	19NE2	74.75
230	25C22	24N	30.81	230	25C22	26N	69.15
230	25C22	24C	47.40	230	25C22	25C	76.13
230	25C22	26CD1	41.43	25CA		19NE2	54.59
25CA	25C22	24N	60.09	25CA	25C22 1		70.00
25CA	25C22	26N	30.49	25CA		24C	30.68
25CA	25C22	25C	15.62	25CA	25C22 1		89.97
25CA	25C22	26CD1	62.21	1510	25C22 1		66.24
1610	25C22	162CA	34.26	19NE2	25C22	24N	44.06

			TA	BLE XIV			
19NE2			75.31	19NE2	25C22	26N	82.03
19NE2		24C	49.24	19NE2	25C22	25C	69.71
19NE2		26CD1	96.23	24N	25C22	26N	68.47
24N	25C22	24C	31.27	24N	25C22	25C	66.84
24N	25C22	26CD1	59.64	162ND1	25C22	26N	88.16
162ND1	25C22	24C	97.87	162ND1	25C22	25C	75.88
162ND1	25C22	162CA	38.60	26N	25C22	24C	39.22
26N	25C22	25C	15.72	26N	25C22	162CA	89.81
26N	25C22	26CD1	37.42	24C	25C22	25C	35.57
24C	25C22	26CD1	47.97	25C	25C22	162CA	86.23
25C	25C22	26CD1	51.39	25SG	25023	25N	74.67
25SG	25023	25CB	38.15	25SG	25023	25CA	54.63
25SG	25023	24C	85.79	25SG	25023	26N	43.64
25SG	25023	25C	46.38	25SG	25023	190E1	92.56
25SG	25023	26CD1	68.54	25SG	25023	162ND1	54.77
25SG	25023	240	79.70	23C	25023	25N	73.70
23C	25023	23CA	31.55	23C	25023	230	24.72
23C	25023	24N	24.08	23C	25023	19NE2	78.98
23C	25023	25CA	97.79	23C	25023	24C	59.01
23C	25023	24CA	36.52	23C	25023	23N	33.47
23C 23C	25023	220	48.06	23C	25023	19CD	90.90
23C	25023	26N	92.04	23C	25023	25C	96.56
23C	25023 25023	26CD1	66.16	23C	25023	22C	38.64
25C 25N	25023 25023	240 2500	64.99	25N	25023	230	75.80
25N	25023	25CB 19NE2	50.26	25N	25023	24N	53.76
25N	25023	19NE2 24C	61.26	25N	25023	25CA	24.14
25N	25023	24C 23N	14.83	25N	25023	24CA	37.18
25N	25023	23N 19CD	97.04	25N	25023	220	79.60
25N	25023	25C	63.80	25N	25023	26N	37.49
25N	25023	26CD1	28.32	25N	25023	190E1	68.00
25N		162ND1	60.48 89.51	25N	25023	22C	87.54
23CA	25023	230	48.80	25N	25023	240	9.64
23CA	25023	19NE2	77.35	23CA 23CA	25023	24N	46.69
23CA	25023	24CA	64.88	23CA 23CA	25023 25023	24C	86.89
23CA	25023	220	38.18	23CA	25023	23N	7.95
23CA	25023	26CD1	93.73	23CA	25023	19CD	87.87
23CA	25023	240	93.16	23CA 230	25023 25023	22C	23.48
230	25023	25CA	97.01	230	25023	24N	42.48
230	25023	24CA	43.63	230	25023	24C	61.75
230	25023	220	72.77	230	25023	23N 26N	53.76
230	25023	25C	89.58	230	25023	26CD1	79.20
230	25023	22C	62.49	230	25023	240	44.93
25CB	25023	19NE2	75.63	25CB	25023	25CA	66.16 26.15
	*			- 3 - 2		~~~~	20.13

			TAI	BLE XIV			
25CB	25023	24C	64.94	25CB	25023	24CA	87.43
25CB	25023	19CD	67.69	25CB	25023	26N	43.69
25CB	25023	25C	31.03	25CB	25023	190E1	57.73
25CB	25023	26CD1	83.37	25CB	25023	162ND1	43.20
25CB	25023	240	59.04	24N	25023	19NE2	59.02
24N	25023	25CA	77.52	24N	25023	24C	40.38
24N	25023	24CA	19.70	24N	25023	23N	44.10
24N	25023	220	40.11	24N	25023	19CD	70.43
24N	25023	26N	81.70	24N	25023	25C	80.42
24N	25023	190E1	85.06	24N	25023	26CD1	70.58
24N	25023	22C	39.18	24N	25023	240	46.65
19NE2	25023	25CA	66.95	19NE2	25023	24C	63.26
19NE2	25023	24CA	65.23	19NE2	25023	23N	69.48
19NE2	25023	220	39.17	19NE2	25023	19CD	11.94
19NE2	25023	26N	96.41	19NE2	25023	25C	81.46
19NE2	25023	190E1	27.37	19NE2	25023	22C	53.87
19NE2	25023	162ND1	76.87	19NE2	25023	240	65.05
25CA	25023	24C	38.79	25CA	25023	24CA	61.29
25CA	25023	220	96.77	25CA	25023	19CD	63.98
25CA	25023	2 <b>6N</b>	31.96	25CA	25023	25C	15.29
25CA	25023	190E1	61.11	25CA	25023	26CD1	68.99
25CA	25023	162ND1	67.28	25CA	25023	240	32.93
24C	25023	24CA	22.55	24C	25023	23N	84.41
24C	25023	220	71.79	24C	25023	19CD	68.91
24C	25023	26N	44.17	24C	25023	25C	40.07
24C	25023	190E1	76.59	24C	25023	26CD1	54.93
24C	25023	22C	77.02	24C	25023	240	6.27
24CA	25023	23N	63.29	24CA	25023	220	57.79
24CA	25023	19CD	74.68	24CA	25023	26N	62.01
24CA	25023	25C	61.72	24CA	25023	190E1	86.76
24CA	25023	26CD1	55.98	24CA	25023	22C	58.75
24CA	25023	240	28.67	23N	25023	220	30.31
23N	25023	19CD	79.92	23N	25023	190E1	93.90
23N	25023	26CD1	98.39	23N	25023	22C	15.63
23N	25023	240	90.66	220	25023	19CD	49.86
220	25023	190E1	64.43	220	25023	22C	14.70
220	25023	240	77.07	19CD	25023	26N	95.38
19CD	25023	25C	79.18	19CD	25023	190E1	15.44
19CD	25023	22C	64.49	19CD	25023		64.94
19CD	25023	240	69.44	26N	25023	25C	16.77
26N	25023	190E1	92.94	26N	25023	26CD1	39.94
26N	25023	162ND1	84.95	26N	25023	240	38.78
25C	25023	190E1	76.17	25C	25023	26CD1	55.41
25C	25023	162ND1	74.20	25C	25023	240	33.82

			ТА	BLE XIV			
190E1	25023	22C	78.82	190E1	25023	162ND1	49.50
190E1	25023	240	75.54	26CD1	25023	240	54.16
22C	25023	240	82.96	162ND1	25023	240	99.10
640	25C24	610D1	49.69	640	25C25	610D1	65.86
640	25C25	64C	4.04	640	25C25	65CA	36.03
640	25C25	61CG	60.48	640	25C25	65N	18.55
610D1	25C25	64C	66.57	610D1	25C25	65CA	65.57
610D1	25C25	61CG	12.80	610D1	25C25	65N	65.27
64C	25C25	65CA	32.21	64C	25C25	61CG	62.08
64C	25C25	65N	14.69	65CA	25C25	61CG	68.81
65CA	25C25	65N	17.52	61CG	25C25	65N	64.30
640	25C26	610D1	75.97	640	25C26	65CA	46.13
640	25C26	65C	60.97	640	25C26	64C	10.65
640	25C26	6 <b>6N</b>	77.86	640	25C26	61CG	67.54
640	25C26	65N	27.99	640	25C26	650	58.42
640	25C26	61CB	50.52	610D1	25C26	65CA	81.67
610D1	25C26	65C	66.87	610D1	25C26	64C	74.15
610D1	25C26	6 <b>6N</b>	75.37	610D1	25C26	61CG	10.88
610D1	25C26	65N	75.81	610D1	25C26	650	51.81
610D1	25C26	61CB	25.47	65CA	25C26	65C	22.26
65CA	25C26	64C	35.83	65CA	25C26	6 <b>6N</b>	33.99
65CA	25C26	61CG	80.84	65CA	25C26	65 <b>N</b>	18.46
65CA	25C26	650	32.38	65CA	25C26	61CB	65.67
65C	25C26	64C	50.43	65C	25C26	6 <b>6N</b>	17.99
65C	25C26	61CG	69.56	65C	25C26	65N	34.42
65C	25C26	650	15.34	65C	25C26	61CB	58.96
64C	25C26	66N	67.22	64C	25C26	61CG	67.20
64C	25C26	65N	17.48	64C	25C26	650	48.92
64C	25C26	61CB	49.45	6 <b>6N</b>	25C26	61CG	80.82
66N	25C26	65N	50.23	6 <b>6</b> N	25C26	650	29.57
66N	25C26	61CB	73.71	61CG	25C26	65N	71.80
61CG	25C26	650	54.22	61CG	25C26	61CB	17.81
65N	25C26	650	36.92	65N	25C26	61CB	54.59
650	25C26	61CB	44.43	610D1	25C27	640	56.92
610D1	25C27	6 <b>6N</b>	69.33	610D1	25C27	65C	57.67
610D1	25C27	65CA	66.06	610D1	25C27	61CG	5.76
640	25C27	66N	63.73	640	25C27	65C	48.11
640	25C27	65CA	35.11	640	25C27	61CG	54.18
66N	25C27	67CE2	75.89	66N	25C27	65C	16.74
66N	25C27	65CA	30.26	66N	25C27	61CG	73.19
66N	25C27	670H	95.43	67CE2	25C27	65C	91.10
67CE2	25C27	670H	29.04	65C	25C27	65CA	18.79
65C	25C27	61CG	60.50	65CA	25C27	61CG	67.06
610D1	25C28	67CE2	87.37	670H	25C28	67CE2	29.20

			TA	BLE XIV			
67CE2	25C30	66CA	87.59	67CE2	25C30	67CD2	20.16
67CE2	25C30	67CZ	18.73	67CE2	25C30	670H	34.56
67CE2	25C30	66C	70.31	67CE2	25C30	660	70.72
67CE2	25C30	67CG	25.18	66N	25C30	610D1	77.24
66N	25C30	66CA	21.80	66N	25C30	67CD2	87.09
66N	25C30	65C	18.44	66N	25C30	66C	34.83
66N	25C30	660	37.41	66N	25C30	650	28.55
66N	25C30	65CA	30.87	66N	25C30	67CG	79.90
610D1	25C30	66CA	79.21	610D1	25C30	65C	61.19
610D1	25C30	66C	99.06	610D1	25C30	650	48.69
610D1	25C30	65CA	65.35	66CA	25C30	67CD2	68.10
66CA	25C30	67CZ	95.64	66CA	25C30	65C	34.85
66CA	25C30	66C	20.50	66CA	25C30	660	32.31
66CA	25C30	650	36.26	66CA	25C30	65CA	51.45
66CA	25C30	67CG	63.14	67CD2	25C30	67CZ	35.10
67CD2	25C30	670H	53.38	67CD2	25C30	66C	52.50
67CD2	25C30	660	56.67	67CD2	25C30	67CG	11.66
67CZ	25C30	670H	19.01	67CZ	25C30	6 <b>6</b> C	75.82
67CZ	25C30	660	70.92	67CZ	25C30	67CG	34.09
65C	25C30	66C	52.09	65C	25C30	660	55.84
65C	25C30	650	14.76	65C	25C30	65CA	18.69
65C	25C30	67CG	97.10	670H	25C30	66C	93.31
670H	25C30	660	85.90	670H	25C30	67CG	53.10
66C	25C30	660	15.93	66C	25C30	650	56.33
66C	25C30	65CA	65.39	66C	25C30	67CG	45.14
660	25C30	650	64.17	660	25C30	65CA	64.29
660	25C30	67CG	46.61	650	25C30	65CA	29.99
650	25C30	67CG	98.82	67CE2	25031	67CZ	25.44
67CE2	25031	670H	43.21	67CE2	25031	67CD2	21.71
67CE2	25031	660	86.23	67CE2	25031	66C	79.23
67CE2	25031	66CA	90.96	67CE2	25031	67CE1	36.09
67CE2	25031	67CG	31.92	67CE2	25031	67CD1	36.21
67CE2	25031	67N	65.41	67CZ	25031	670H	23.53
67CZ	25031	67CD2	40.63	67CZ	25031	660	90.28
67CZ	25031	66C	90.73	67CZ	25031	67CE1	17.83
67CZ	25031	67CG	39.73	67CZ	25031	67CD1	30.41
67CZ	25031	67N	78.58	670H	25031	67CD2	62.56
670H	25031	67CE1	36.02	670H	25031	67CG	63.25
670н	25031	67CD1	52.26	67CD2	25031	660	66.16
67CD2	25031	66N	90.96	67CD2	25031	66C	57.61
67CD2	25031	66CA	69.91	67CD2	25031	67CE1	41.91
67CD2	25031	67CG	16.99	57CD2	25031	67CD1	31.58
67CD2	25031	67N	43.75	660	25031	66N	42.55
660	25031	66C	18.09	660	25031	66CA	35.81

			TA	BLE XIV			
660	25031	67CE1	75.32	660	25031	67CG	54.80
660	25031	65C	57.68	660	25031		59.89
660	25031	67N	27.36	660	25031		66.84
66N	25031	66C	37.22	66N	25031	66CA	21.24
66N	25031	67CG	88.11	66N	25031	65C	15.16
66N	25031	67CD1	99.35	66N	25031	67N	49.30
66N	25031	65CA	28.53	66C	25031	66CA	21.98
66C	25031	67CE1	79.50	66C	25031	67CG	51.32
66C	25031	65C	51.46	66C	25031	67CD1	62.32
66C	25031	67N	13.91	66C	25031	65CA	65.42
66CA	25031	67CE1	99.90	66CA	25031	67CG	68.85
66CA	25031	65C	32.23	66CA	25031	67CD1	82.55
66CA	25031	67N	30.44	66CA	25031	65CA	48.87
67CE1	25031	67CG	33.08	67CE1	25031	67CD1	17.35
67CE1	25031	67N	69.51	67CG	25031	67CD1	17.79
67CG	25031	67N	38.86	65C	25031	67N	62.38
65C	25031	65CA	17.93	67CD1	25031	67N	52.17
67N	25031	65CA	77.82	660	25C32	66N	41.03
660	25C32	67CZ	76.35	660	25C32	670H	95.33
660	25C32	67CE2	69.48	660	25C32	66C	15.10
660	25C32	66CA	31.92	660	25C32	65CA	69.81
660	25C32	65C	56.36	660	25C32	67CE1	66.82
660	25C32	67CD2	53.70	6 <b>6N</b>	25C32	67CE2	87.72
66N	25C32	66C	33.67	66N	25C32	66CA	17.88
66N	25C32	65CA	31.90	6 <b>6N</b>	25C32	65C	15.39
66N	25C32	67CD2	73.68	67CZ	25C32	670H	19.19
67CZ	25C32	67CE2	19.23	67CZ	25C32	66C	73.63
67CZ	25C32	66CA	85.66	67CZ	25C32	67CE1	16.63
67CZ	25C32	67CD2	30.02	670H	25C32	67CE2	33.49
670H	25C32	66C	92.69	670 <b>H</b>	25C32	67CE1	30.65
670H	25C32	67CD2	47.69	67CE2	25C32	66C	62.32
67CE2	25C32	66CA	70.20	67CE2	25C32	65C	96.55
67CE2	25C32	67CE1	30.87	67CE2	25C32	67CD2	16.03
66C	25C32	66CA	19.46	66C	25C32	65CA	65.30
66C	25C32	65C	48.62	6 <b>6C</b>	25C32	67CE1	68.20
66C	25C32	67CD2	46.37	66CA	25C32	65CA	49.19
66CA	25C32	65C	30.83	6GCA	25C32	67CE1	84.10
66CA	25C32	67CD2	55.84	65CA	25C32	65C	19.14
65C	25C32	67CD2	84.41	67CE1	25C32	67CD2	33.97
670H	25033	67CZ	16.87	670H	25033	66N	97.69
670H	25033	67CE2	28.97	67 <b>0H</b>		1600	89.11
67CZ	25033	66N	81.70	67CZ	25033	67CE2	16.30
67CZ		1600	93.56	66N	25033	67CE2	69.22
660	25C34	6 <b>6</b> N	39.13	650	25C34	65CA	71.17

			TA	BLE XIV			
660	25C34	25 <i>S</i> G	87.16	660	25C34	65C	53.66
660	25C34	66C	9.91	66N	25C34	65CA	32.59
66N	25C34	25 <i>S</i> G	85.90	66N	25C34	65C	14.62
66N	25C34	66C	30.25	1610	25C34	161C	15.14
1610	25C34	25SG	60.22	1610	25C34	1600	61.47
65CA	25C34	25 <b>S</b> G	79.10	65CA	25C34	65C	18.14
65CA	25C34	66C	62.76	161C	25C34	25 <i>S</i> G	68.39
161C	25C34	1600	48.09	25 <i>S</i> G	25C34	65C	84.17
25 <i>S</i> G	25C34	66C	90.62	65C	25C34	66C	44.86
1610	25C35	1600	72.48	1610	25C35	161C	18.16
1610	25C3 <del>5</del>	162N	31.48	1610	25C35	161CA	33.51
1610	25C35	163N	64.70	1610	25C35	160C	64.27
1610	25C35	25SG	63.17	1610	25C35	162CA	35.94
1610	25C35	161N	48.76	1610	25C35	162C	54.37
1600	25C35	161C	58.42	1600	25C35	162N	63.10
1600	25C35	161CA	39.04	1600	25C35	160C	13.43
1600	25C35	162CA	80.67	1600	25C35	161N	27.44
1600	25C35	162C	86.34	161C	25C35	162N	17.93
161C	25C35	161CA	20.68	161C	25C35	163N	61.49
161C	25C35	160C	48.25	161C	25C35	25 <i>S</i> G	75.24
161C	25C35	162CA	31.05	161C	25C35	161N	32.29
161C	25C35	162C	47.54	660	25C35	25 <i>S</i> G	85.01
162N	25Ç35	161CA	32.91	162N	25C35	163N	45.91
162N	25C35	160C	50.42	162N	25C35	25SG	71.74
162N	25C35	162CA	17.88	162N	25C35	161N	35.87
162N	25C35	162C	30.79	161CA	25C35	163N	78.80
161CA	25C35	160C	31.32	161CA	25C35	25 <i>S</i> G	95.35
161CA	25C35	162CA	49.73	161CA	25C35	161N	17.33
161CA	25C35	162C	63.45	1631	25C35	160C	88.14
163N	25C35	25 <i>S</i> G	47.65	163N	25C35	162CA	30.56
163N	25C35	161N	77.99	163N	25C35	162C	15.86
160C	25C35	162CA	67.69	160C	25C35	161N	15.98
160C	25C35	162C	72.94	25 <b>S</b> G	25C35	162CA	55.42
25 <i>S</i> G	25C35		57.22	162CA	25C35	161N	53.66
162CA	25C35	162C	18.43	161N	25C35	162C	62.17
660	25C36	163CB	99.95	660	25C36	66C	0.80
660	25C36	25 <b>S</b> G	91.26	660	25C36	68SD	70.79
660	25C36	25CB	44.37	163CB	25C36		35.50
163CB	25C36	163CA	19.65	163CB	25C36	134CB	72.59
163CB	25C36	162C	49.56	163CB	25C36	25SG	58.45
163CB	25C36	68SD	45.87	163CB	25C36		79.29
163CB	25C36	161C	92.40	163CB	25C36	26CB	57.89
163N	25C36	163CA	19.24	163N	25C36		72.07
163N	25C36	162C	15.65	163N	25C36	25 <i>S</i> G	48.74

			TA	BLE XIV			
163N	25C36	1600	94.75	163N	25C36	68SD	80.23
163N	25C36	162N	44.07	163N	25C36	161C	56.92
163N	25C36	26CB	84.19	163CA	25C36	134CB	63.35
163CA	25C36	162C	30.81	163CA	25C36	25SG	59.31
163CA	25C36	68SD	61.48	163CA	25C36	162N	60.75
163CA	25C36	161C	74.90	163CA	25C36	26CB	76.21
134CB	25C36	162C	66.02	134CB	25C36	1600	71.49
134CB	25C36	68SD	70.15	134CB	25C36	162N	74.57
134CB	25C36	161C	86.37	162C	25C36	25SG	57.47
162C	25C36	1600	79.13	162C	25C36	68SD	91.74
162C	25C36	162N	29.94	162C	25C36	161C	44.33
162C	25C36	26CB	99.68	66C	25C36	25 <b>S</b> G	91.85
66C	25C36	68SD	71.28	66C	25C36	26CB	45.17
25 <i>S</i> G	25C36	68SD	94.27	25SG	25C36	162N	65.87
25 <b>S</b> G	25C36	161C	65.98	25 <b>S</b> G	25C36	26CB	58.75
1600	25C36	162N	52.97	1600	25C36	161C	46.61
68SD	25C36	26CB	51.40	162N	25C36	161C	15.56
660	25C37	67CD1	72.85	660	25C37	67CE1	84.55
660	25C37	66C	7.48	660	25C37	67CG	55.90
660	25C37	67CZ	77.27	660	25C37	67CA	36.66
660	25C37	68SD	75.55	660	25C37	163CB	87.96
660	25C37	68CE	96.44	660	25C37	67N	20.80
67CD1	25C37	67CE1	21.00	67CD1	25C37	209CD2	54.83
67CD1	25C37	66C	65.41	67CD1	25C37	67CG	16.96
67CD1	25C37	67CZ	32.06	67CD1	25C37	67CA	45.91
67CD1	25C37	68SD	86.51	67CD1	25C37	68CE	88.92
67CD1	25C37	67N	53.96	67CE1	25C37	209CD2	57.96
67CE1	25C37	66C	77.24	67CE1	25C37	67CG	32.77
67CE1	25C37	67CZ	16.58	67CE1	25C37	67CA	64.93
67CE1	25C37	67 <b>N</b>	69.10	209CD2	25C37	134CB	54.20
209CD2	25C37	67CG	69.67	209CD2	25C37	67CZ	74.22
209CD2	25C37	67CA	82.39	20 <b>9CD2</b>	25C37	68SD	72.35
209CD2	25C37	68CE	57.34	209CD2	25C37	67N	98.27
209CD2	25C37	1600	99.57	6 <b>6</b> C	25C37	67CG	48.46
66C	25C37	67CZ	70.63	6 <b>6C</b>	25C37	67CA	31.49
66C	25C37	68SD	76.55	66C	25C37	163CB	93.59
66C	25C37	68CE	96.76	6 <b>6</b> C	25C37	67N	14.69
134CB	25C37	68SD	73.29	134CB	25C37		66.42
134CB	25C37	68CE	52.90	134CB	25C37	1600	70.87
67CG	25C37	67CZ	36.23	67 <b>CG</b>	25C37	67CA	32.41
67CG	25C37	68SD	83.13	67 <b>CG</b>	25C37	68CE	91.57
67CG	25C37	67N	37.47	67CZ	25C37	67CA	67.51
67CZ	25C37	67N	66.38	67CZ	25C37		95.75
67CA	25C37	68SD	57.53	67CA	25C37	163CB	92.22

		TA	BLE XIV		
67CA	25C37 68CE	73.28	67CA	25C37 67N	17.05
68SD	25C37 163CB	43.37	68SD	25C37 68CE	21.25
68SD	25C37 67N	68.73	163CB	25C37 68CE	46.72
163CB	25C37 67N	95.05	68CE	25C37 67N	87.13
660	25C38 26CB	65.13	660	25C38 66C	6.46
660	25C38 26CA	86.69	660	25C38 26N	98.55
660	25C38 68SD	86.25	660	25C38 26CG	59.16
660	25C38 26CD1	64.07	660	25C38 66N	34.26
660	25C38 67CA	29.20	660	25C38 67N	14.41
660	25C38 66CA	18.99	163CB	25C38 26CB	81.15
163CB	25C38 25SG	73.30	163CB	25C38 163N	37.84
163CB	25C38 26CA	60.80	163CB	25C38 26N	60.85
163CB	25C38 163CA	18.86	163CB	25C38 68SD	54.22
163CB	25C38 26CG	95.41	163CB	25C38 162C	48.20
26CB	25C38 25SG	80.09	26CB	25C38 66C	58.81
26CB	25C38 26CA	21.77	26CB	25C38 26N	35.93
26CB	25C38 163CA	98.31	26CB	25C38 68SD	66.22
26CB	25C38 26CG	18.06	26CB	25C38 26CD1	34.61
26CB	25C38 66N	65.47	26CB	25C38 67CA	61.54
26CB	25C38 67N	56.77	26CB	25C38 66CA	58.47
25SG	25C38 163N	55.98	25SG	25C38 26CA	68.25
25SG	25C38 26N	47.87	25SG	25C38 163CA	68.75
25SG	25C38 26CG	73.64	25\$G	25C38 26CD1	61.85
25SG	25C38 66N	91.11	25SG	25C38 162C	60.63
163N	25C38 26CA	87.46	163N	25C38 26N	76.37
163N	25C38 163CA	20.53	163N	25C38 68SD	89.44
163N	25C38 162C	12.27	66C	25C38 26CA	80.45
66C	25C38 26N	92.09	66C	25C38 68SD	84.88
66C	25C38 26CG	52.75	66C	25C38 26CD1	58.32
66C	25C38 66N	32.59	66C	25C38 67CA	29.60
66C	25C38 67N	12.87	66C	25C38 66CA	15.91
26CA	25C38 26N	20.39	26CA	25C38 163CA	77.00
26CA	25C38 68SD	62.23	26CA	25C38 26CG	34.72
26CA	25C38 26CD1	45.79	26CA	25C38 66N	84.74
26CA	25C38 67CA	78.68	26CA	25C38 67N	77.24
26CA	25C38 66CA	79.85	26CA	25C38 162C	99.66
26N	25C38 163CA	72.27	26N	25C38 68SD	79.71
26N	25C38 26CG	39.78	26N	25C38 26CD1	42.14
26N	25C38 66N	85.55	26N	25C38 67CA	97.07
26N	25C38 67N	92.29	26N	25C38 66CA	86.55
26N	25C38 162C	87.67	163CA	25C38 68SD	68.96
163CA	25C38 162C	29.60	68SD	25C38 26CG	83.61
68SD	25C38 67CA	57.57	6850	25C38 67N	72.28
68SD	25C38 66CA	98.54	6 <b>8S</b> D	25C38 162C	96.25

			TA	BLE XIV			
26CG	25C38	26CD1	17.49	26CG	25C38	66N	50.02
26CG	25C38	67CA	65.74	26CG	25C38	67N	55.48
26CG	25C38	66CA	47.14	26CD1	25C38	66N	43.47
26CD1	25C38	67CA	78.32	26CD1	25C38	67N	64.92
26CD1	25C38	66CA	47.51	6 <b>6N</b>	25C38	67CA	62.18
66N	25C38	67N	45.30	66N	25C38	66CA	17.02
67CA	25C38	67N	17.03	67CA	25C38	66CA	45.35
67N	25C38	66CA	28.38	65CA	25C39	6 <b>6</b> N	38.23
65CA	25C39	660	77.80	65CA	25C39	65C	20.22
65CA	25C39	26CD1	57.59	65CA	25C39	65N	12.86
65CA	25C39	230	51.79	65CA	25C39	66CA	49.23
25SG	25C39	1610	71.00	25SG	25C39	26CD1	64.51
25SG	25C39	161C	75.18	25SG	25C39	230	59.23
66N	25C39	660	39.58	66N	25C39	65C	18.09
66N	25C39	26CD1	47.02	66N	25C39	65N	50.53
66N	25C39	230	71.29	66N	25C39	66CA	11.02
660	25C39	65C	57.60	660	25C39	26CD1	57.39
660	25C39	65N	89.89	660	25C39	230	96.66
660	25C39	66CA	28.57	1610	25C39	161C	12.79
65C	25C39	26CD1	48.36	65C	25C39	65N	32.45
65C	25C39	230	59.69	65C	25C39	66CA	29.05
26CD1	25C39	65N	60.85	26CD1	25C39	230	40.56
26CD1	25C39	66CA	47.30	65N	25C39	230	44.53
65N	25C39	66CA	61.43	230	25C39	66CA	77.82
66N	25040	65CA	49.46	66N	25040	26CD1	65.76
66N	25040	65C	23.27	66N	25040	6 <b>6</b> 0	49.28
66N	25040	66CA	14.40	6 <b>6N</b>	25040	26CG	63.42
66N	25040	65 <b>N</b>	61.07	66N	25040	26NE1	58.78
66N	25040	66C	34.93	6 <b>6N</b>	25040	230	92.25
66N	25040	26CB	74.00	6 <b>6N</b>	25040	650	24.57
65CA	25040	26CD1	79.08	65CA	25040	65C	27.14
65CA	25040	660	98.43	65CA	25040	66CA	63.57
65CA	25040	26CG	90.14	65CA	25040	65N	13.53
65CA	25040	26NE1	63.17	65CA	25040	66C	84.39
65CA	25040	230	63.91	65CA	25040	650	28.53
26CD1	25040	65C	65.35	26CD1	25040	25SG	80.06
26CD1	25040	660	75.51	26CD1	25040	66CA	61.87
26CD1	25040	26CG	17.43	26CD1	25040	65N	76.84
26CD1	25040	26 <b>NE</b> 1	16.20	26CD1	25040	66C	66.92
25CD1	25040	230	50.44	26CD1	25040	26CB	36.73
26CD1	25040	650	60.57	26CD1	25040	26N	42.40
65C	25040	660	72.49	65C	25040	66CA	36.77
65C	25040	26CG	70.55	65 <b>C</b>	25040	65N	37.88
65C	25040	26NE1	52.57	65C	25040	66C	57.85

			TA	BLE XIV			
65C	25040	230	74.40	65C	25040	26CB	86.99
65C	25040	650	4.93	25SG	25040	26CG	82.32
25SG	25040	26NE1	89.52	25SG	25040	230	68.61
25SG	25040	26CB	75.26	25SG	25040	26N	44.90
660	25040	66CA	36.06	660	25040	26CG	60.15
660	25040	26NE1	82.09	660	25040	66C	15.20
660	25040	26CB	51.97	660	25040	650	73.07
660	25040	26N	81.84	66CA	25040	26CG	55.19
66CA	25040	65N	74.60	66CA	25040	26NE1	59.26
66CA	25040	66C	21.11	66CA	25040	230	98.28
66CA	25040	26CB	62.10	66CA	25040	650	37.02
66CA	25040	26N	91.86	26CG	25040	65N	90.75
26CG	25040	26NE1	31.46	26CG	25040	66C	54.19
26CG	25040	230	67.47	26CG	25040	26CB	20.67
26CG	25040	650	66.40	26CG	25040	26N	38.44
65N	25040	26NE1	60.66	65N	25040	66C	95.71
65N	25040	230	52.18	65N	25040	650	37.84
26NE1	25040	66C	70.37	26NE1	25040	230	42.57
26NE1	25040	26CB	51.93	26NE1	25040	650	47.64
26NE1	25040	26N	57.31	66C	25040	26CB	52.36
66C	25040	650	58.10	66C	25040	26N	83.81
230	25040	26CB	83.39	230	25040	650	70.59
230	25040	26N	67.33	26CB	25040	650	83.51
26CB	25040	26N	31.46	25SG	25N41	1610	86.79
25 <i>S</i> G	25N41	161C	87.19	25 <b>S</b> G	25N41	230	62.77
25 <i>S</i> G	25 <b>N41</b>	6 <b>6N</b>	93.69	1610	25 <b>N4</b> 1	161C	11.62
65CA	25N41	230	51.79	65CA	25N41	65N	15.59
65CA	25N41	6 <b>6N</b>	29.14	230	25N41	65N	44.44
230	25N41	6 <b>6N</b>	64.51	6 <b>5N</b>	25N41	66N	44.25
25 <i>S</i> G	25N42	230	85.08	25 <b>S</b> G	25N42	23CA	98.91
25 <i>S</i> G	25N42	23C	83.87	25SG	25N42	1610	78.66
25SG	25N42	25CB	15.26	25 <b>SG</b>	25N42	25N	39.74
25 <b>S</b> G	25N42	26CD1	63.71	25 <b>S</b> G	25N42	24N	72.85
230	25 <b>N4</b> 2	23CA	37.86	230	25N42	23C	18.24
230	25 <b>N4</b> 2	65CA	60.94	230	25N42	25CB	79.36
230	25N42	65N	52.05	230	25 <b>N4</b> 2	25N	48.33
230	25N42	26CD1	43.36	230	25N42	24N	25.15
23CA	25N42	23C	22.71	23CA	25N42	65CA	87.12
23CA	25N42	25CB	85.89	23CA	25N42	65N	70.74
23CA	25N42	25N	60.69	23CA	25N42	26CD1	80.64
23CA	25N42	24N	29.02	23C	25N42	65CA	77.61
23C	25N42	25CB	74.03	23C	25N42	65N	65.78
23C	25N42	25N	44.16	23C	25N42	26CD1	58.52
23C	25N42	24N	11.74	65CA	25N42	65N	18.58

			TA	BLE XIV			
65CA	25N42	25 <b>N</b>	98.97	65CA	25N42	26CD1	51.99
65CA	25N42	24N	86.08	1610	25N42	25CB	84.39
25CB	25N42	25N	31.12	25CB	25N42	26CD1	68.91
25CB	25N42	24N	62.41	65N	25N42	25N	97.62
65N	25N42	26CD1	59.56	65N	25N42	24N	76.12
25N	25N42	26CD1	50.78	25N	25N42	24N	33.18
26CD1	25N42	24N	57.77	660	25N43	66N	50.46
660	25 <b>N4</b> 3	66C	15.63	660	25N43	66CA	35.78
660	25N43	65CA	83.97	660	25N43	65C	66.03
660	25N43	67CZ	75.44	660	25N43	67CE2	67.55
660	25 <b>N4</b> 3	67CE1	68.04	660	25N43	67 <b>N</b>	16.30
66N	25N43	66C	39.54	66N	25N43	66CA	19.49
66N	25N43	65CA	35.70	66N	25N43	65C	15.67
66N	25 <b>N4</b> 3	67CE2	84.94	66N	25N43	67N	47.36
66C	25 <b>N4</b> 3	66CA	21.89	66C	25N43	65CA	75.04
66C	25N43	65C	55.08	66C	25N43	67CZ	72.67
66C	25 <b>N4</b> 3	67CE2	61.03	66C	25N43	67CE1	69.65
66C	25 <b>N4</b> 3	67 <b>N</b>	8.63	6 <b>6CA</b>	25N43	65CA	54.94
66CA	25N43	65C	34.13	66CA	25N43	67CZ	83.41
66CA	25 <b>N4</b> 3	67CE2	68.23	66CA	25 <b>N43</b>	67CE1	85.39
66CA	25N43	67N	28.69	65CA	25N43	65C	21.08
65CA	25N43	67N	83.04	65C	25N43	67CE2	93.39
65C	25N43	67N	62.65	67CZ	25N43	67CE2	16.72
67CZ	25N43	67CE1	16.22	67CZ	25N43	67N	64.53
67CE2	25 <b>N4</b> 3	67CE1	28.45	67CE2	25N43	67N	53.83
67CE1	25N43	67N	61.03				

TABLE XV

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
184CD1	25C1	184CB	39.64	184CD1	25C1	184CG	20.72
184CD1	25C1	1840	79.67	184CD1	25C1	180D1	92.08
184CD1	25C1	184CA	45.35	184CD1	25C1	184C	65.97
184CD1	25C1	18CG	95.35	184CD1	25C1	200	95.11
184CD1	25C1	184NE1	15.46	184CD1	25C1	184CD2	28.37
184CB	25C1	184CG	22.78	184CB	25C1	1840	43.71
184CB	25C1	180D1	91.44	184CB	25C1	184CA	22.80
184CB	25C1	18ND2	86.39	184CB	25C1	184C	35.31
184CB	25C1	18CG	84.04	184CB	25C1	184NE1	51.55
184CB	25C1	184CD2	34.26	184CG	25C1	1840	66.36
184CG	25C1	184CA	38.32	184CG	25C1	184C	56.18
184CG	25C1	18CG	98.73	184CG	25C1	184NE1	29.35
184CG	25C1	184CD2	15.76	1840	25C1	180D1	69.22
1840	25C1	184CA	35.76	1840	25C1	18ND2	48.03
1840	25C1	184C	16.21	1840	25C1	18CG	54.71
1840	25C1	184NE1	93.86	1840	25C1	184CD2	77.07
180D1	25C1	184CA	68.65	180D1	25C1	18ND2	32.37
180D1	25C1	184C	63.21	180D1	25C1	18CG	16.11
180D1	25C1	200	62.46	184CA	25C1	18ND2	67.26
184CA	25C1	184C	20.72	184CA	25C1	18CG	62.00
184CA	· 25C1	184NE1	60.54	184CA	25C1	184CD2	53.33
18ND2	25C1	184C	51.16	18ND2	25C1	18CG	17.75
18ND2	25C1	200	89.83	184C	25C1	18CG	51.58
184C	25C1	184NE1	80.98	184C	25C1	184CD2	69.47
18CG	25C1	200	77.96	200	25C1	184NE1	90.38
184NE1	25C1	184CD2	28.12	180 <b>D1</b>	25C2	184CA	82.66
180D1	25C2	200	82.02	180D1	25C2	18CG	17.46
180D1	25C2	20N	42.19	180D1	25C2	18ND2	35.11
180D1	25C2	1840	74.48	180D1	25C2	19CG	79.73
180D1	25C2	184C	70.99	180 <b>D1</b>	25C2	1830	67.05

			TA	BLE XV			
180D1	25C2	20CA	48.76	180D1	25C2	20C	68.51
180D1	25C2	19N	41.49	180D1	25C2	184N	80.79
180D1	25C2	19C	47.40	180D1	25C2	183C	74.42
180D1	25C2	18CB	20.67	184CD1	25C2	184CA	49.69
184CD1	25C2	184CG	20.47	184CD1	25C2	184CB	40.02
184CD1	25C2	1840	80.18	184CD1	25C2	19CG	63.56
184CD1	25C2	184C	69.76	184CD1	25C2	184NE1	15.99
184CD1	25C2	1830	53.74	184CD1	25C2	19N	85.37
184CD1	25C2	184N	44.38	184CD1	25C2	184CD2	23.21
184CD1	25C2	183C	46.49	184CA	25C2	18CG	70.97
184CA	25C2	184CG	39.55	184CA	25C2	18ND2	72.06
184CA	25C2	184CB	22.96	184CA	25C2	1840	35.00
184CA	25C2	19CG	83.11	184CA	25C2	184C	20.65
184CA	25C2	184NE1	65.57	184CA	25C2	1830	40.23
184CA	25C2	19N	70.98	184CA	25C2	184N	13.04
184CA	25C2	184CD2	51.91	184CA	25C2	183C	28.28
184CA	25C2	18CB	62.55	200	25C2	18CG	98.28
200	25C2	20N	45.78	200	25C2	19CG	69.54
200	25C2	20CA	33.26	200	25C2	20C	13.52
200	25C2	19N	80.94	200	25C2	2410H2	94.87
200	25C2	19C	50.12	18CG	25C2	20N	59.60
18CG	25C2	18ND2	19.96	18CG	25C2	184CB	91.12
18CG	25C2	1840	57.44	18CG	25C2	19CG	91.35
18CG	25C2	184C	56.05	18CG	25C2	1830	66.77
18CG	25C2	20CA	65.32	18CG	25C2	20C	84.82
18CG	25C2	19N	51.19	18CG	25C2	18 <b>4N</b>	72.42
18CG	25C2	19C	64.04	18CG	25C2	183C	70.48
18CG	25C2	18CB	11.35	184CG	25C2	184CB	22.17
184CG	25C2	1840	63.20	184CG	25C2	19CG	82.47
184CG	25C2	184C	56.51	184CG	25C2	184NE1	31.65
184CG	25C2	1830	60.77	184CG	25C2	19N	95.86
184CG	25C2	184N	40.46	184CG	25C2	184CD2	12.76
184CG	25C2	183C	49.79	20 <b>N</b>	25C2	18ND2	76.27
20N	25C2	19CG	52.54	20 <b>N</b>	25C2	1830	72.94
20N	25C2	20CA	19.39	20 <b>N</b>	25C2	20C	34.03
20N	25C2	19N	38.18	20 <b>N</b>	25C2	184N	99.39
20N	25C2	19C	11.58	20N	25C2	183C	85.38
20N	25C2	1.8CB	59.41	19ND2	25C2	184CB	87.16

			TA	ABLE XV			
18ND2	25C2	1840	47.43	18ND2	25C2	184C	52.81
18ND2	25C2	1830	80.93	18ND2	25C2	20CA	77.11
18ND2	25C2	20C	94.81	18ND2	25C2	19N	70.92
18ND2	25C2	184N	77.97	18ND2	25C2	2410H2	87.90
18ND2	25C2	19C	82.49	18ND2	25C2	183C	81.11
18ND2	25C2	18CB	28.59	184CB	25C2	1840	41.08
184CB	25C2	19CG	93.42	184CB	25C2	184C	35.36
184CB	25C2	184NE1	53.47	184CB	25C2	1830	58.06
184CB	25C2	19N	92.03	184CB	25C2	184N	30.81
184CB	25C2	2410H2	98.67	184CB	25C2	184CD2	32.23
184CB	25C2	183C	45.37	184CB	25C2	18CB	84.11
1840	25C2	184C	16.82	1840	25C2	184NE1	94.48
1840	25C2	1830	69.02	1840	25C2	19N	87.53
1840	25C2	184N	47.05	1840	25C2	2410H2	86.44
1840	25C2	184CD2	71.78	1840	25C2	183C	59.90
1840	25C2	18CB	55.54	19CG	25C2	184C	97.61
19CG	25C2	184NE1	62.79	19CG	25C2	1830	45.36
19CG	25C2	20CA	67.99	19CG	25C2	20C	68.52
19CG	25C2	19N	40.19	19CG	25C2	184N	70.06
19CG	25C2	19C	41.14	19CG	25C2	184CD2	86.47
19CG	25C2	183C	55.31	19CG	25C2	18CB	82.88
184C	25C2	184NE1	85.37	184C	25C2	1830	52.27
184C	25C2	19N	73.95	184C	25C2	184N	31.18
184C	25C2	184CD2	67.55	184C	25C2	183C	43.16
184C	25C2	18CB	50.40	184NE1	25C2	1830	65.82
184NE1	25C2	19N	93.08	184NE1	25 <b>C2</b>	184N	60.22
184NE1	25C2	184CD2	27.19	184NE1	25C2	183C	60.67
1830	25C2	20CA	92.25	1830	25C2	19N	35.35
1830	25C2	184N	28.06	1830	25C2	19C	63.84
1830	25C2	184CD2	71.98	1830	25C2	183C	12.70
1830	25C2	18CB	55.43	20CA	25C2	20C	19.77
20CA	25C2	19N	57.18	20CA	25C2	19C	29.86
20CA	25C2	18CB	69.00	20C	25C2	19N	71.24
20C	25C2	2410H2	99.96	20C	25C2	19C	40.54
20C	25C2	18CB	88.76	19N	25C2	184N	61.22
19N	25C2	19C	30.91	19N	25C2	183C	47.43
19N	25C2	18CB	42.94	134N	25C2	19C	91.32
184N	25C2	184CD2	53.18	184N	25C2	183C	15.60

			TA	ABLE XV			
184N	25C2	18CB	62.40	2410H2	25C2	184CD2	94.01
19C	25C2	183C	76.51	19C	25C2	18CB	61.35
184CD2	25C2	183C	61.73	183C	25C2	18CB	59.34
200	25C3	19CG	91.80	200	25C3	20N	53.46
200	25C3	180D1	84.67	200	25C3	20C	13.71
200	25C3	20CA	35.80	200	25C3	19CD	97.07
200	25C3	19C	58.10	200	25C3	19N	89.89
200	25C3	19CB	82.19	200	25C3	18CG	93.78
200	25C3	19CA	76.43	200	25C3	19NE2	84.73
200.	25C3	21N	7.78	184CD1	25C3	19CG	71.43
184CD1	25C3	184NE1	19.86	184CD1	25C3	19CD	64.18
184CD1	25C3	184CG	15.82	184CD1	25C3	1830	49.75
184CD1	25C3	19N	83.17	184CD1	25C3	184CA	39.91
184CD1	25C3	19CB	82.78	184CD1	25C3	18CG	94.02
184CD1	25C3	19CA	92.64	184CD1	25C3	19NE2	76.30
184CD1	25C3	184CB	30.78	184CD1	25C3	190E1	50.83
184CD1	25C3	184CE2	22.67	19CG	25C3	20N	61.01
19CG	25C3	180D1	81.71	19CG	25C3	20C	85.09
19CG	25C3	184NE1	72.26	19CG	25C3	20CA	79.11
19CG	25C3	19CD	19.52	19CG	25C3	184CG	85.19
19CG	25C3	19C	46.68	19CG	25C3	1830	45.20
19CG	25C3	19N	40.95	19CG	25C3	184CA	79.09
19CG	25C3	19CB	14.28	19CG	25C3	18CG	86.69
19CG	25C3	19CA	32.54	19CG	25C3	19NE2	29.74
19CG	25C3	184CB	89.69	19CG	25C3	190E1	27.37
19CG	25C3	21N	86.08	19CG	25C3	184CE2	83.99
20N	25C3	180D1	41.77	20N	25C3	20C	39.98
20N	25C3	20CA	21.31	2 <b>0N</b>	25C3	19CD	78.54
20N	25C3	19C	14.86	20 <b>N</b>	25C3	1830	71.47
20 <b>N</b>	25C3	19N	37.66	20 <b>N</b>	25C3	184CA	95.61
20N	25C3	19CB	46.74	20 <b>N</b>	25C3	18CG	52.23
20N	25C3	19CA	30.40	20 <b>N</b>	25C3	19NE2	78.92
20N	25C3	190E1	88.38	20 <b>N</b>	25C3	21N	45.77
180D1	25C3	20C	71.90	18 <b>0D1</b>	25C3	20CA	50.00
180D1	25C3	184CG	95.90	180 <b>D1</b>	25C3	19C	51.61
180D1	25C3	1830	58.74	130D1	25C3	19N	40.87
180D1	25C3	184CA	63,.14	180D1	25C3	19CB	70.42
180D1	25C3	18CG	10.53	180D1	25C3	19CA	52.41

			TA	ABLE XV			
180D1	25C3	184CB	79.52	180D1	25C3	21N	78.64
20C	25C3	20CA	22.29	20C	25C3	19CD	94.40
20C	25C3	19C	46.10	20C	25C3	19N	77.00
20C	25C3	19CB	73.57	20C	25C3	18CG	81.47
20C	25C3	19CA	64.92	20C	25C3	19NE2	84.78
20C	25C3	21N	6.74	184NE1	25C3	19CD	58.87
184NE1	25C3	184CG	31.08	184NE1	25C3	1830	64.98
184NE1	25C3	19N	95.87	184NE1	25C3	184CA	59.73
184NE1	25C3	19CB	85.90	184NE1	25C3	19NE2	66.72
184NE1	25C3	184CB	48.57	184NE1	25C3	190E1	46.24
184NE1	25C3	184CE2	11.83	20CA	25C3	19CD	94.19
20CA	25C3	19C	33.03	20CA	25C3	1830	92.18
20CA	25C3	19N	58.66	20CA	25C3	19CB	65.28
20CA	25C3	18CG	59.95	20CA	25C3	19CA	51.05
20CA	25C3	19NE2	90.04	20CA	25C3	21N	28.97
19CD	25C3	184CG	79.74	19CD	25C3	19C	63.74
19CD	25C3	1830	55.90	19CD	25C3	19N	60.04
19CD	25C3	184CA	84.73	19CD	25C3	19CB	32.93
19CD	25C3	19CA	51.74	19CD	25C3	19NE2	15.91
19CD	25C3	184CB	89.68	19CD	25C3	190E1	13.39
19CD	25C3	21N	93.33	19CD	25C3	184CE2	70.59
184CG	25C3	1830	54.91	184CG	25C3	19N	88.52
184CG	25C3	184CA	32.83	184CG	25C3	19CB	95.04
184CG	25C3	18CG	87.10	184CG	25C3	19NE2	92.12
184CG	25C3	184CB	17.79	184CG	25C3	190E1	66.46
184CG	25C3	184CE2	27.27	19C	25C3	1830	65.15
19C	25C3	19N	32.44	19C	25 <b>C</b> 3	184CA	94.94
19C	25C3	19CB	32.53	19C	25C3	18CG	61.42
19C	25C3	19CA	18.94	19C	25 <b>C</b> 3	19NE2	64.39
19C	25C3	190E1	73.97	19C	25C3	21N	50.41
1830	25C3	19N	33.87	1830	25C3	184CA	35.08
1830	25C3	19CB	47.01	1830	25C3	18CG	56.73
1830	25C3	19CA	47.13	1830	25C3	19NE2	71.34
1830	25C3	184CB	50.06	1830	25C3	190E1	50.42
1830	25C3	184CE2	72.00	19K	25C3	184CA	62.75
19N	25C3	19CB	31.18	19N	25C3	1.8CG	46.02
19N	25C3	19CA	17.94	19N	25C3	19NE2	69.90
19N	25C3	184CB	80.68	19N	25C3	190E1	63.48

			TA	ABLE XV			
19N	25C3	21N	82.11	184CA	25C3	19CB	82.09
184CA	25C3	18CG	54.80	184CA	25C3	19CA	79.48
184CA	25C3	184CB	18.58	184CA	25C3	190E1	74.55
184CA	25C3	184CE2	59.43	19CB	25C3	18CG	76.90
19CB	25C3	19CA	18.91	19CB	25C3	19NE2	39.32
19CB	25C3	184CB	95.96	19CB	25C3	190E1	41.65
19CB	25C3	21N	75.62	19CB	25C3	184CE2	97.49
18CG	25C3	19CA	59.97	18CG	25C3	184CB	70.15
18CG	25C3	21N	88.18	19CA	25C3	19NE2	57.54
19CA	25C3	184CB	96.61	19CA	25C3	190E1	59.34
19CA	25C3	21N	68.88	19NE2	25C3	190E1	26.89
19NE2	25C3	21N	82.33	19NE2	25C3	184CE2	77.62
184CB	25C3	190E1	77.28	184CB	25C3	184CE2	44.87
190E1	25C3	184CE2	58.06	200	25C4	20C	5.29
200	25C4	19CG	73.30	200	25C4	20N	37.40
200	25C4	20CA	22.70	200	25C4	19CD	84.35
200	25C4	180D1	61.35	200	25C4	19NE2	78.63
184CD1	25C4	184NE1	19.87	184CD1	25C4	19CG	58.10
184CD1	25C4	20 <b>N</b>	90.23	184CD1	25C4	184CG	14.72
184CD1	25C4	19CD	55.97	184CD1	25C4	184CE2	27.01
184CD1	25C4	180D1	75.63	184CD1	25C4	19NE2	69.64
184NE1	25C4	19CG	63.14	184NE1	25C4	184CG	28.26
184NE1	25C4	19CD	53.91	184NE1	25C4	184CE2	14.23
184NE1	25C4	180D1	94.59	184NE1	25C4	19NE2	63.81
20C	25C4	19CG	69.98	20C	25C4	20N	32.25
20C	25C4	20CA	17.47	20C	25C4	19CD	82.18
20C	25C4	180D1	56.22	20C	25C4	19NE2	77.77
19CG	25C4	20 <b>N</b>	46.51	19CG	25C4	184CG	71.25
19CG	25C4	20CA	62.49	19CG	25C4	19CD	18.45
19CG	25C4	184CE2	77.31	19CG	25C4	180D1	59.85
19CG	25C4	19NE2	29.19	20 <b>N</b>	25C4	184CG	96.60
20N	25C4	20CA	17.63	20 <b>N</b>	25C4	19CD	63.68
20N	25C4	180D1	30.95	20N	25C4	19NE2	67.31
184CG	25C4	19CD	70.61	184CG	25C4	184CE2	27.62
184CG	25C4	180D1	75.70	184CG	25C4	2410H2	87.84
184CG	25C4	19NE2	84.36	20CA	25C4	19CD	78.25
20CA	25C4	180D1	38.85	20CA	25C4	2410H2	97.61
20CA	25C4	19NE2	78.54	19CD	25C4	184CE2	67.82

			TA	BLE XV			
19CD	25C4	180D1	78.00	19CD	25C4	19NE2	15.45
184CE2	25C4	2410H2	99.63	184CE2	25C4	19NE2	76.63
180D1	25C4	2410H2	82.91	180D1	25C4	19NE2	87.95
200	25C5	184CD1	97.18	184CD1	25C5	184NE1	18.35
184CD1	25C5	184CG	16.64	184CD1	25C5	184CE2	27.73
184CD1	25C5	184CD2	26.39	184NE1	25C5	184CG	28.44
184NE1	25C5	184CE2	16.30	184NE1	25C5	184CD2	26.54
184CG	25C5	184CE2	27.87	184CG	25C5	184CD2	16.56
184CE2	25C5	184CD2	16.34	184CD1	25C6	184CG	18.54
184CD1	25C6	184CB	33.62	184CD1	25C6	184NE1	17.14
184CD1	25C6	200	88.08	184CD1	25C6	184CD2	28.33
184CD1	25C6	184CE2	26.95	184CG	25C6	184CB	19.13
184CG	25C6	184NE1	29.00	184CG	25C6	184CD2	17.43
184CG	25C6	184CE2	27.74	184CB	25C6	184NE1	47.54
184CB	25C6	184CD2	32.76	184CB	25C6	184CE2	46.38
184NE1	25C6	200	88.62	184NE1	25C6	184CD2	27.82
184NE1	25C6	184CE2	16.20	184CD2	25C6	184CE2	16.68
200	25C7	20C	5.94	200	25C7	19NE2	91.02
200	25C7	19CG	72.12	200	25C7	19CD	89.31
200	25C7	21CA	34.14	200	25C7	220	65.12
200	25C7	21N	17.69	200	25C7	21C	46.47
20C	25C7	19NE2	86.13	20C	25C7	19CG	69.33
20C	25C7	19CD	85.63	20C	25C7	21CA	31.54
20C	25C7	220	59.23	20C	25C7	21N	14.16
20C	25C7	21C	42.01	19NE2	25 <b>C</b> 7	19CG	32.17
19NE2	25C7	184NE1	66.44	19NE2	25C7	19CD	17.05
19NE2	25C7	21CA	93.37	19NE2	25 <b>C7</b>	220	37.10
19NE2	25C7	184CD1	69.09	19NE2	25C7	21N	87.99
19NE2	25C7	21C	79.11	19CG	25C7	184NE1	58.99
19CG	25C7	19CD	19.56	19CG	25C7	21CA	91.19
19CG	25C7	220	50.36	19CG	25 <b>C7</b>	184CD1	52.04
19CG	25C7	21N	77.89	19CG	25C7	21C	84.98
184NE1	25C7	19CD	53.40	184NE1	25 <b>C7</b>	184CD1	16.66
19CD	25C7	220	49.22	19CD	25C7	184CD1	53.17
19CD	25C7	21N	91.50	19CD	25C7	21C	90.15
21CA	25C7	220	56.87	21CA	25 <b>C7</b>	21N	17.50
21CA	25C7	21C	18.03	220	25C7	21N	55.41
220	25C7	210	42.02	21N	25C7	21C	28.94

			T.	ABLE XV			
19NE2	2508	19CD	20.48	19NE2	2508	184NE1	81.39
19NE2	2508	200	89.30	19NE2	2508	19CG	35.60
19NE2	2508	190E1	30.68	19NE2	2508	220	40.85
19NE2	2508	184CD1	79.20	19NE2	2508	184CE2	92.69
19CD	2508	184NE1	63.10	19CD	2508	200	86.55
19CD	2508	19CG	21.39	19CD	2508	190E1	15.46
19CD	2508	220	55.33	19CD	2508	184CD1	58.99
19CD	2508	184CE2	75.44	184NE1	2508	19CG	65.39
184NE1	2508	190E1	50.75	184NE1	2508	184CD1	16.57
184NE1	2508	184CE2	13.19	200	2508	19CG	66.59
200	2508	190E1	99.26	200	2508	220	62.20
200	2508	184CD1	90.68	19CG	2508	190E1	32.68
19CG	2508	220	53.93	19CG	2508	184CD1	54.99
19CG	2508	184CE2	78.58	190E1	2508	220	69.68
190E1	2508	184CD1	50.54	190E1	2508	184CE2	62.13
184CD1	2508	184CE2	26.99	19NE2	25C9	184NE1	68.02
19NE2	25C9	19CD	15.81	184NE1	25C9	19CD	52.53
162ND1	25C11	184CZ2	58.70	162ND1	25C11	162CE1	15.57
184CZ2	25C11	162CE1	49.86	162ND1	25C14	162CG	20.80
162ND1	25C14	162CB	38.90	162ND1	25C14	162CE1	17.70
162ND1	25C14	162CA	43.05	162ND1	25C14	1610	71.97
162ND1	25C14	184CZ2	66.72	162ND1	25C14	162CD2	25.28
162ND1	25C14	1610D1	87.80	162ND1	25C14	162N	60.63
162ND1	25C14	162NE2	23.20	162ND1	25C14	161C	71.88
162ND1	25C14	25SG	43.89	162CG	25C14	162CB	21.88
162CG	25C14	162CE1	32.41	162CG	25C14	162CA	35.32
162CG	25C14	1610	70.65	162 <b>CG</b>	25C14	184CZ2	61.27
162CG	25C14	162CD2	14.29	162CG	25C14	1610D1	69.93
162CG	25C14	162N	50.52	162CG	25C14	162NE2	26.85
162CG	25C14	161C	65.13	162CG	25C14	25SG	60.65
162CB	25C14	162CE1	53.68	162CB	25C14	162CA	20.45
162CB	25C14	1610	55.44	162CB	25C14	184CZ2	75.70
162CB	25C14	162CD2	33.63	162CB	25C14	1610D1	49.02
162CB	25C14	162N	30.67	162CB	25C14	162NE2	48.52
162CB	25C14	161C	46.38	162CB	25C14	25SG	66.67
162CE1	25C14	162CA	60.62	162CE1	25C14	1610	88.83
162CE1	25C14	184CZ2	55.07	162CE1	25C14	162CD2	27.92
162CE1	25C14	162N	78.11	162CE1	25C14	162NE2	14.38

		T	ABLE XV		
162CE1	25C14 161C	89.56	162CE1	25C14 25SG	51.14
162CA	25C14 1610	35.99	162CA	25C14 184CZ2	94.99
162CA	25C14 162CI	2 49.43	162CA	25C14 1610D1	50.42
162CA	25C14 162N	17.63	162CA	25C14 162NE2	60.81
162CA	25C14 161C	30.12	162CA	25C14 25SG	53.36
1610	25C14 162CD	84.93	1610	25C14 1610D1	54.15
1610	25C14 162N	27.79	1610	25C14 162NE2	93.77
1610	25C14 161C	14.72	1610	25C14 25SG	53.31
184CZ2	25C14 162CD	2 47.18	184CZ2	25C14 1610D1	98.34
184CZ2	25C14 162NE	2 43.71	162CD2	25C14 1610D1	77.69
162CD2	25C14 162N	63.80	162CD2	25C14 162NE2	16.69
162CD2	25C14 161C	78.91	162CD2	25C14 25SG	69.06
1610D1	25C14 162N	36.11	1610D1	25C14 162NE2	94.34
1610D1	25C14 161C	39.94	162N	25C14 162NE2	77.07
162N	25C14 161C	15.92	162N	25C14 25SG	64.86
162NE2	25C14 161C	90.92	162NE2	25C14 25SG	64.04
161C	25C14 25SG	63.90	162CB	25015 162ND1	47.26
162CB	25015 162CG	27.43	162CB	25015 1610D1	63.34
162CB	25015 162CA	23.06	162CB	25015 162CE1	60.10
162CB	25015 162CD	2 40.37	162CB	25015 162N	35.92
162CB	25015 184CZ	2 91.76	162CB	25015 162NE2	55.17
162CB	25015 161C	52.53	162CB	25015 1610	60.45
162CB	25015 137CB	54.06	162CB	25015 161CG	69.17
162CB	25015 1370	87.23	162CB	25015 161CB	73.81
162CB	25015 184CH	2 88.25	162ND1	25015 162CG	25.12
162ND1	25015 162CA	49.83	162ND1	25015 162CE1	16.98
162ND1	25015 162CD	2 32.89	162ND1	25015 162N	70.32
162ND1	25015 184CZ	2 74.09	162ND1	25015 162NE2	26.96
162ND1	25015 161C			25015 1610	74.28
162ND1	25015 137CB	88.45	162ND1	25015 184CH2	81.61
162CG	25015 1610D	90.25	162CG	25015 162CA	41.57
				25015 162CD2	
162CG	25015 162N	60.35	162CG	25015 184CZ2	70.97
162CG	25015 162NE2	28.08	162CG	25015 161C	74.63
162CG	25015 1610	77.02	162CG	25015 137CB	63.77
162CG	25015 161CG	96.60	162CG	25015 1370	97.22
162CG	25015 184CH2	72.59	151001	25015 162CA	61.52
1610D1	25015 162CD2	98.29	161001	25015 162N	43.32

			TA	ABLE XV			
1610D1	25015	161C	45.80	1610D1	25015	1610	60.74
1610D1	25015	137CB	55.90	1610D1	25015	161CG	11.21
1610D1	25015	1370	59.03	1610D1	25015	161CB	28.61
162CA	25015	162CE1	66.36	162CA	25015	162CD2	58.17
162CA	25015	162N	20.56	162CA	25015	162NE2	68.37
162CA	25015	161C	33.06	162CA	25015	1610	37.86
162CA	25015	137CB	73.81	162CA	25015	161CG	62.82
162CA	25015	161CB	60.45	162CE1	25015	162CD2	31.95
162CE1	25015	162N	86.92	162CE1	25015	184CZ2	59.12
162CE1	25015	162NE2	17.11	162CE1	25015	161C	95.94
162CE1	25015	1610	90.42	162CE1	25015	137CB	90.07
162CE1	25015	184CH2	68.68	162CD2	25015	162N	75.70
162CD2	25015	184CZ2	53.94	162CD2	25015	162NE2	17.99
162CD2	25015	161C	91.05	162CD2	25015	1610	94.26
162CD2	25015	137CB	58.26	162CD2	25015	1370	88.29
162CD2	25015	184CH2	55.47	162N	25015	162NE2	88.07
162N	25015	161C	17.35	162N	25015	1610	30.00
162N	25015	137CB	72.58	162N	25015	161CG	42.77
162N	25015	1370	95.02	162N	25015	161CB	40.21
184CZ2	25015	162NE2	47.88	184CZ2	25015	137CB	69.04
184CZ2	25015	1370	71.73	184CZ2	25015	184CH2	14.31
162NE2	25015	1610	99.69	162NE2	25015	137CB	74.12
162NE2	25015	184CH2	54.65	161C	25015	1610	15.92
161C	25015	137CB	87.40	161C	25015	161CG	40.49
161C	25015	161CB	30.37	1610	25015	161CG	53.91
1610	25015	181CB	39.98	137CB	25015	161CG	67.10
137CB	25015	1370	34.26	137CB	25015	161CB	84.07
137CB	25015	184CH2	<b>56</b> .32	161CG	25015	1370	67.49
161CG	25015	161CB	17.89	1370	25015	161CB	84.60
1370	25015	184CH2	57.69	162ND1	25N16	25 <i>S</i> G	57.00
162ND1	25N16	1610	81.16	162ND1	25N16	162CE1	17.28
162ND1	25พ16	162CG	17.24	162ND1	25N16	162CA	44.41
162ND1	25N16	162CB	35.55	162ND1	25N15	161C	76.08
162ND1	25N16	25CB	47.59	162ND1	25N16	162N	61.35
162ND1	25N16	19NE2	78.00	25 <b>SG</b>	25N16	1610	66.73
25SG	25N16	162CE1	60.44	25 <b>SG</b>	25N16	162CG	69.41
25 <i>S</i> G	25N16	162CA	62.55	25SG	25N16	162CB	74.95
25SG	25N16	161C	75.51	25SG	25N16	25CB	22.32

	TABLE XV							
25SG	25N16 162N	74.12	25 <i>S</i> G	25N16 19NE2	65.05			
1610	25N16 162CE1	97.79	1610,	25N16 162CG	74.22			
1610	25N16 162CA	38.96	1610	25N16 162CB	57.51			
1610	25N16 161C	14.31	1610	25N16 25CB	86.73			
1610	25N16 162N	28.02	162CE1	25N16 162CG	31.36			
162CE1	25N16 162CA	61.69	162CE1	25N16 162CB	51.34			
162CE1	25N16 161C	93.35	162CE1	25N16 25CB	44.14			
162CE1	25N16 162N	78.58	162CE1	25N16 19NE2	62.97			
162CG	25N16 162CA	35.27	162CG	25N16 162CB	20.22			
162CG	25N16 161C	65.76	162CG	25N16 25CB	63.85			
162CG	25N16 162N	49.75	162CG	25N16 19NE2	94.18			
162CA	25N16 162CB	20.52	162CA	25N16 161C	31.69			
162CA	25N16 25CB	71.57	162CA	25N16 162N	17.61			
162CB	25N16 161C	47.01	162CB	25N16 25CB	76.68			
162CB	25N16 162N	30.74	161C	25N16 25CB	92.93			
161C	25N16 162N	16.27	25CB	25N16 162N	87.03			
25CB	25N16 19NE2	47.50	25SG	25N17 162ND1	74.24			
25SG	25N17 162CA	87.90	25SG	25N17 162CG	86.46			
25SG	25N17 162CB	97.30	25SG	25N17 25CB	24.15			
25 <b>S</b> G	25N17 162CE1	70.19	25 <b>S</b> G	25N17 163N	54.13			
25SG	25N17 162C	71.67	25 <b>S</b> G	25N17 162CD2	84.79			
1610	25N17 162CA	52.13	1610	25N17 161C	16.50			
1610	25N17 162CG	92.62	1610	25N17 162CB	71.30			
1610	25N17 162N	34.71	1610	25N17 163N	65.25			
1610	25N17 162C	55.77	1610	25N17 161CA	17.53			
162ND1	25N17 162CA	56.42	162ND1	25N17 161C	95.91			
162ND1	25N17 162CG	18.78	162ND1	25N17 162CB	41.07			
162ND1	25N17 162N	76.00	162ND1	25N17 25CB	57.57			
162ND1	25N17 162CE1	14.79	162ND1	25N17 163N	54.56			
162ND1	25N17 162C	55.20	162ND1	25N17 162CD2	12.12			
162CA	25N17 161C	40.29	162CA	25N17 162CG	41.09			
162CA	25N17 162CB	23.23	162CA	25N17 162N	21.53			
162CA	25N17 25CB	91.56	162CA	25N17 162CE1	71.02			
162CA	25N17 163N	33.82	162CA	25N17 162C	16.39			
162CA	25N17 161CA	50.27	162CA	25N17 162CD2	49.51			
161C	25N17 162CG	78.57	161C					
161C	25N17 162N	19.96	161C					
161C	25N17 162C	48.62	161C	25N17 161CA	10.96			

			TA	BLE XV			
161C	25N17	162CD2	87.05	162CG	25N17	162CB	22.57
162CG	25N17	162N	58.64	162CG	25N17	25CB	73.95
162CG	25 <b>N1</b> 7	162CE1	33.00	162CG	25N17	163N	51.02
162CG	25 <b>N</b> 17	162C	44.96	162CG	25N17	161CA	86.42
162CG	25 <b>N1</b> 7	162CD2	8.55	162CB	25N17	162N	36.69
162CB	25 <b>N1</b> 7	25CB	91.50	162CB	25N17	162CE1	55.54
162CB	25N17	163N	48.31	162CB	25N17	162C	34.62
162CB	25N17	161CA	63.87	162CB	25N17	162CD2	30.83
162N	25N17	162CE1	90.79	162N	25N17	163N	50.48
162N	25 <b>N1</b> 7	162C	33.85	162N	25N17	161CA	29.03
162N	25 <b>N1</b> 7	162CD2	67.15	25CB	25N17	162CE1	49.52
25CB	25 <b>N1</b> 7	163N	61.60	25CB	25N17	162C	77.64
25CB	25N17	162CD2	69.50	162CE1	25N17	163N	64.51
162CE1	25 <b>N17</b>	162C	68.30	162CE1	25N17	162CD2	25.14
163N	25 <b>N</b> 17	162C	17.55	163N	25N17	161CA	73.19
163N	25 <b>N17</b>	162CD2	56.08	162C	25N17	161CA	59.51
162C	25N17	162CD2	52.20	161CA	25N17	162CD2	94.68
184NE1	25C18	19NE2	72.19	184NE1	25C18	184CZ2	34.15
184NE1	25C18	162ND1	64.76	184NE1	25C18	162CE1	49.26
184NE1	25C18	19CD	55.65	184NE1	25C18	190E1	48.22
184NE1	25C18	184CE2	16.77	19NE2	25C18	162ND1	74.92
19NE2	25C18	162CE1	65.60	19NE2	25C18	19CD	16.66
19NE2	25C18	190E1	29.19	19NE2	25C18	184CE2	88.46
184CZ2	25C18	162ND1	61.13	184CZ2	25C18	162CE1	54.08
184CZ2	25C18	19CD	87.16	184CZ2	25C18	190E1	75.81
184CZ2	25C18	184CE2	17.74	162ND1	25C18	162CE1	17.06
162ND1	25C18	19CD	67.56	162ND1	25C18	190E1	52.95
162ND1	25C18	184CE2	64.20	162CE1	25C18	19CD	54.80
162CE1	25C18	190E1	39.41	162CE1	25C18	184CE2	52.06
19CD	25C18	190E1	15.45	19CD	25C18	184CE2	71.81
190E1	25C18	184CE2	62.64	25 <b>S</b> G	25C19	25CB	30.92
25SG	25C19	1610	96.60	25SG	25C19	162ND1	68.37
25SG	25C19	25N	49.94	25 <b>5G</b>	25C19	25CA	32.83
25SG	25C19	230	90.89	25 <b>SG</b>	25C19	162CE1	65.52
25SG	25C19	162CA	73.52	25 <b>S</b> G	25C19	19NE2	86.96
25 <i>S</i> G	25C19	23C	92.63	25 <b>5</b> G	25C19		42.91
25SG	25C19	161C	94.€4	25 <b>SG</b>	25C19	162CG	71.57
25SG	25C19	190E1	68.21	25 <b>S</b> G	25C19	25C	20.12

		TA	ABLE XV		
25SG	25C19 26N	28.60	25CB	25C19 162N	D1 59.73
25CB	25C19 25N	36.66	25CB	25C19 25C	A 17.74
25CB	25C19 230	86.02	25CB	25C19 162C	E1 48.75
25CB	25C19 162CA	86.55	25CB	25C19 19N	E2 56.95
25CB	25C19 23C	79.82	25CB	25C19 163N	62.99
25CB	25C19 23CA	90.53	25CB	25C19 162C	G 68.32
25CB	25C19 19OE1	37.70	25CB	25C19 25C	26.97
25CB	25C19 26N	42.42	1610	25C19 162N	D1 80.02
1610	25C19 162CE1	96.21	1610	25C19 162C	A 38.78
1610	25C19 163N	57.58	1610	25C19 161C	9.12
1610	25C19 162CG	68.74	162ND1	25C19 25N	95.15
162ND1	25C19 25CA	77.47	162ND1	25C19 162C	E1 16.21
162ND1	25C19 162CA	42.69	162ND1	25C19 19N	E2 79.46
162ND1	25C19 163N	49.40	162ND1	25C19 161C	71.09
162ND1	25C19 162CG	11.30	162ND1	25C19 190	E1 54.54
162ND1	25C19 25C	81.37	162ND1	25C19 26N	95.47
25N	25C19 25CA	20.26	25N	25C19 230	49.36
25N	25C19 162CE1	81.63	25N	25C19 19N	E2 50.17
25N	25C19 23C	44.73	25N	25C19 163N	92.43
25N	25C19 23CA	59.48	25N	25C19 190	E1 52.21
25N	25C19 25C	30.33	25N	25C19 26N	33.58
25CA	25C19 23O	69.09	25CA	25C19 162C	E1 66.04
25CA	25C19 19NE2	56.58	25CA	25C19 23C	64.91
25CA	25C19 163N	73.45	25CA	25C19 23C	A 78.58
25CA	25C19 162CG	85.89	25 <b>CA</b>	25C19 190	E1 46.47
25CA	25C19 25C	17.28	25 <b>CA</b>	25C19 26N	29.70
230	25C19 19NE2	67.81	230	25C19 23C	15.74
230	25C19 23CA	30.61	230	25C19 190	E1 88.93
230	25C19 25C	71.99	230	25C19 26N	63.05
162CE1	25C19 162CA	58.48	162CE1	25C19 19N	E2 63.81
162CE1	25C19 163N	59.21	162CE1	25C19 161C	87.29
162CE1	25C19 162CG	27.50	162CE1	25C19 190	E1 38.34
162CE1	25C19 25C	73.76	162CE1	25C19 26N	89.08
1.62CA	25C19 163N	31.32	162CA	25C19 161C	30.71
162CA	25C19 162CG	31.93	162CA	25C19 190	E1 96.39
162CA	25C19 25C	93.49	162CA	25C19 26N	99.54
19NE2	25C19 23C	52.36	19NE2	25C19 23C	A 47.51
19NE2	25C19 162CG	90.36	19NE2	25C19 190	E1 27.54

TABLE XV							
19NE2	25C19	25C	73.76	19NE2	25C19	26N	82.78
23C	25C19	23CA	18.96	23C	25C19	190E1	75.13
23C	25C19	25C	72.57	23C	25C19	26N	67.65
163N	25C19	161C	53.30	163N	25C19	162CG	44.32
163N	25C19	190E1	88.05	163N	25C19	25C	63.02
163N	25C19	26N	68.23	23CA	25C19	190E1	74.37
23CA	25C19	25C	89.19	23CA	25C19	26N	86.09
161C	25C19	162CG	59.79	162CG	25C19	190E1	65.83
162CG	25C19	25C	87.18	162CG	25C19	26N	99.91
190E1	25C19	25C	62.33	190E1	25C19	26N	76.16
25C	25C19	26N	15.58	25SG	25020	25CB	34.29
25 <b>S</b> G	25020	25N	58.41	25 <b>SG</b>	25020	230	94.11
25 <i>S</i> G	25020	162ND1	57.05	25 <i>S</i> G	25020	25CA	42.72
25SG	25020	190E1	79.03	25 <i>S</i> G	25020	19CD	92.36
25SG	25020	24N	95.32	25 <i>S</i> G	25020	162CE1	62.12
25SG	25020	1610	62.41	25 <b>S</b> G	25020	24C	65.93
25CB	25020	19NE2	70.39	25CB	25020	23C	93.15
25CB	25020	25N	40.44	25CB	25020	230	94.44
25CB	25020	162ND1	54.11	25CB	25020	25CA	19.78
25CB	25020	190E1	46.09	25CB	25020	19CD	58.16
25CB	25020	24N	78.39	25CB	25020	162CE1	47.36
25CB	25020	1610	92.37	25CB	25020	24C	49.33
19NE2	25020	23CA	63.45	19NE2	25020	23C	67.92
19NE2	25020	25N	61.15	19NE2	25020	230	85.94
19NE2	25020	162ND1	88.31	19NE2	25020	25CA	65.49
19NE2	25020	190E1	31.85	19NE2	25020	19CD	15.16
19NE2	25020	24N	55.12	19NE2	25020	162CE1	71.33
19NE2	25020	24C	60.20	19NE2	25020	23N	58.44
23CA	25020	23C	24.00	23CA	25020	25N	73.06
23CA	25020	230	37.74	23CA	25020	25CA	93.15
23CA	25020	190E1	94.06	23 <b>CA</b>	25020	19CD	77.95
23CA	25020	24N	33.98	23CA	25020	24C	64.37
23CA	25020	23N	6.16	23C	25020	25N	52.77
23C	25020	230	18.86	23C	25020	25CA	73.50
23C	25020	190E1	92.10	23C	25020	19CD	79.18
23C	25020	24N	16.93	23C	25020	24C	43.85
23C	25020	23N	28.31	25N	25020	230	55.74
25N	25020	162ND1	94.17	25N	25020	25CA	20.73

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			TA	BLE XV			
25N	25020	190E1	59.69	25N	25020	19CD	59.03
25N	25020	24N	39.08	25N	25020	162CE1	84.24
25N	25020	24C	8.93	25N	25020	23N	74.25
230	25020	25CA	75.11	230	25020	19CD	95.79
230	25020	24N	31.46	230	25020	24C	47.72
230	25020	23N	43.41	162ND1	25020	25CA	73.82
162ND1	25020	190E1	58.12	162ND1	25020	19CD	73.91
162ND1	25020	162CE1	17.08	162ND1	25020	1610	63.55
25CA	25020	190E1	51.34	25CA	25020	19CD	57.52
25CA	25020	24N	59.36	25CA	25020	162CE1	65.59
25CA	25020	24C	29.65	25CA	25020	23N	93.68
190E1	25020	19CD	16.69	190E1	25020	24N	75.92
190E1	25020	162CE1	41.10	190E1	25020	24C	64.20
190E1	25020	23N	89.60	19CD	25020	24N	64.32
19CD	25020	162CE1	56.83	19CD	25020	24C	60.79
19CD	25020	23N	73.22	24N	25020	24C	30.42
24N	25020	23N	35.59	162CE1	25020	1610	80.58
162CE1	25020	24C	92.22	24C	25020	23N	65.85
160CD1	25C21	1580	97.34	160CD1	25C21	160CG	23.97
160CD1	25C21	160CB	37.71	160CD1	25C21	160N	66.46
160CD1	25C21	158C	87.61	160CD1	25C21	160CA	53.04
1580	25C21	160CG	75.13	1580	25C21	160CB	77.51
1580	25C21	160N	47.16	1580	25C21	158C	10.74
1580	25021	160CA	63.99	160CG	25C21	160CB	22.63
160CG	25C21	160N	43.47	150CG	25C21	158C	66.36
160CG	25C21	16UCA	33.10	160CB	25C21	160N	33.28
160CB	25C21	158C	71.92	160CB	25C21	160CA	16.60
160N	25C21	158C	44.97	1 E ON	25C21	160CA	17.43
158C	25C21	160CA	60.36	160CD1	25C22	160CG	21.04
160CD1	25C22	160CB	35.29	160CD1	25C22	209CD2	57.75
160CD1	25C22	209CD1	70.22	160CD1	25C22	1580	76.29
160CG	25C22	160CB	20.85	150 <b>CG</b>	25C22	209CD2	73.30
160CG	25C22	209CD1	90.52	2.60CG	25C22	1580	57.61
160CB	25C22	209CD2	69.60	160CB	25C22	209CD1	94.36
160CB	25C22	1580	61.33	209CD2	25C22	209CD1	29.68
160CD1	25C23	209CD2	57.43	160CD1	25C23	160CB	34.09
160CD1	25C23	1600	71.45	160CD1	25C23	160CG	18.98
209CD2	25C23	67CE1	66.23	209CD2	25C23	160CB	73.03

		Т	ABLE XV		
209CD2	25C23 1600	88.66	209CD2	25C23 670	H 97.28
209CD2	25C23 160CG	72.08	209CD2	25C23 67C	- 1 1 2 3
67CE1	25C23 67OH	31.06	67CE1	25C23 67C	
160CB	25C23 1600	37.90	160CB	25C23 160C	
1600	25C23 160CG	56.76	670H	25C23 67C	
1600	25C24 160CB	44.46	1600	25C24 160C	11.63
1600	25C24 160CA	31.06	1600	25C24 160N	41.50
1600	25C24 160CD1	75.56	1600	25C24 160C	
160CB	25C24 160C	34.19	160CB	25C24 160C	. –
160CB	25C24 160N	32.07	160CB	25C24 160CI	
160CB	25C24 160CG	18.03	160C	25C24 160C	
160C	25C24 160N	31.04	160C	25C24 160CI	
160C	25C24 160CG	51.26	160CA	25C24 160N	18.21
160CA	25C24 160CD1	50.13	160CA	25C24 160CG	33.24
160N	25C24 160CD1	57.42	160N	25C24 160CG	38.76
160CD1	25C24 160CG	18.66	67CE1	25C24 67OH	28.53
1600	25C25 160N	52.68	1600	25C25 160CE	48.40
1600	25C25 160CA	37.54	1600	25C25 160C	16.03
1600	25C25 160CG	68.51	1600	25C25 160CD	1 77.48
1600	25C25 159C	58.15	1600	25C25 159CA	73.59
1600	25C25 158C	96.27	160N	25C25 160CB	39.49
160N	25C25 158O	51.80	160N	25C25 160CA	21.57
160N	25C25 160C	36.87	160N	25C25 160CG	45.04
160N	25C25 160CD1	63.96	160N	25C25 159C	10.14
160N	25C25 159CA	27.45	160N	25C25 158C	43.72
160CB	25C25 1580	77.37	160CB	25C25 160CA	22.69
160CB	25C25 160C	37.96	160CB	25C25 160CG	20.22
160CB	25C25 160CD1	31.90	160CB	25C25 159C	49.57
160CB	25C25 159CA	65.68	160CB	25C25 158C	69.44
	25C25 160CA		1580	25C25 160C	88.59
1580	25C25 160CG	65.33	1580	25C25 160CD	77.45
1580		46.06	1580	25C25 159CA	33.23
1580	<del>-</del>		160CA	25C25 160C	22.25
	25C25 160CG		160CA	25C25 160CD	L 53.48
160CA	25C25 159C		160CA	25C25 159CA	48.92
160CA				25C25 160CG	
160C	25C25 160CD1		160C	25C25 159C	43.18
160C	25C25 159CA	59.72	150C	25C25 158C	80.59

			T	ABLE XV			
160CG	25C25	5 160CD1	19.02	160CG	25C25	159C	53.50
160CG	25C25	159CA	65.18	160CG	25C25	158C	58.66
160CD1	25C25	159C	72.53	160CD1	25C25	159CA	83.50
160CD1	25C25	158C	72.33	159C	25C25	159CA	17.81
159C	25C25	158C	38.66	159CA	25C25	158C	27.96
1580	25C26	160N	61.78	1580	25C26	160CB	92.43
1580	25C26	160CG	82.57	1580	25C26	158C	10.10
1580	25C26	160CA	80.14	1580	25C26	160CD1	98.52
1580	25C26	159C	50.29	1580	25C26	159CA	34.65
1580	25C26	160C	92.37	1580	25C26	159N	20.44
1580	25C26	158CA	18.72	160N	25C26	160CB	40.08
160N	25C26	160CG	50,16	160N	25C26	158C	54.73
160N	25C26	160CA	20.14	160N	25C26	160CD1	70.80
160N	25C26	1600	45.75	160N	25C26	159C	12.46
160N	25C26	159CA	31.73	160N	25C26	160C	31.11
160N	25C26	159N	41.35	160N	25C26	158CA	65.61
160CB	25C26	160CG	23.66	160CB	25C26	158C	82.75
160CB	25C26	160CA	21.78	160CB	25C26	160CD1	36.20
160CB	25C26	1600	42.04	160CB	25C26	159C	51.71
160CB	25C26	159CA	70.60	160CB	25C26	160C	32.83
160CB	25C26	159N	74.02	160CB	25C26	158CA	86.07
160CG	25C26	158C	72.51	160CG	25C26	160CA	39.22
160CG	25C26	160CD1	21.40	160CG	25C26	1600	65.66
160CG	25C26	159C	58.12	160CG	25C26	159CA	73.32
160CG	25C26	160C	55.17	160CG	25C26	15 <b>9N</b>	69.08
160CG	25C26	158CA	70.57	158C	25C26	160CA	71.90
158C	25C26	160CD1	88.97	158C	25C26	1600	99.96
158C	25C26	159C	44.20	158C	25C26	159CA	31.81
158C	25C26	160C	85.81	158C	25C26	159N	14.55
158C	25C26	158CA	15.01	160CA	25C26	160CD1	56.75
160CA	25C26	1600	33.81	160CA	25C26	159C	32.58
160CA	25C26	159CA	51.87	160CA	25C26	160C	19.16
160CA	25C26	159N	59.89	150CA	25C26	158CA	80.10
160CD1	25C26	1600	74.72	160CD1	25C26	159C	79.44
160CD1	25C26	159CA	94.53	160CD1	25C26	160C	68.62
160CD1	25C26	159N	88.47	160CD1	25C26	158CA	83.00
1600	25C26	159C	55.78	1600	25C26	159CA	71.36
1600	25C26	160C	15.23	1.600	25C26	159N	85.80

		T.	ABLE XV		
159C	25C26 1590		159C	25C26 160C	42.08
159C	25C26 1591	30.09	159C	25C26 158CA	56.63
159CA	25C26 1600	59.37	159CA	25C26 159N	17.84
159CA	25C26 158C	A 46.52	160C	25C26 159N	72.10
160C	25C26 158C	A 96.40	159N	25C26 158CA	28.80
1600	25C27 160C	5.58	1600	25C27 160CB	36.45
670H	25C27 67C	E1 31.65	670H	25C27 67CZ	17.00
67CE1	25C27 67C	Z 17.03	160C	25C27 160CB	31.24
1600	25028 1600	3.01	1600	25028 161CA	33.47
1600	25028 1600	B 35.39	1600	25028 161N	15.24
1600	25028 160C	A 17.00	1600	25028 1610	59.68
1600	25028 161C	45.54	160C	25028 161CA	33.37
160C	25028 160C	B 34.06	160C	25028 161N	14.62
160C	25028 160C	A 16.50	160C	25028 1610	58.60
160C	25028 161C	44.40	161CA	25028 160CB	65.80
161CA	25028 161N	18.92	161CA	25028 160CA	49.84
161CA	25028 1610	28.96	161CA	25028 161C	17.13
160CB	25028 161N	47.18	160CB	25028 160CA	19.23
160CB	25028 1610	84.88	160CB	25028 161C	71.68
161N	25028 160C	A 30.98	161N	25028 1610	44.50
161N	25028 161C	30.43	160CA	25028 1610	73.79
160CA	25028 161C	59.64	1610	25028 161C	14.21
1600	25C29 160C	8.30	1600	25C29 1610	64.77
1600	25C29 161C	A 35.29	1600	25C29 161C	50.76
1600	25C29 161N	20.86	160C	25C29 1610	58.92
160C	25C29 161C	30.98	160C	25C29 161C	44.50
160C	25C29 161N	14.62	1610	25C29 161CA	30.36
1610	25C29 161C	14.65	1610	25C29 66O	98.33
1610	25C29 161N	44.35	161CA	25C29 161C	18.41
161CA	25C29 161N		1.61C	25C29 161N	30.04
660	25C29 67CE	67.43	67CE1	25030 660	85.21
67CE1	25030 67CI	18.89	67CE1	25030 67CZ	18.55
67CE1	25030 66C	76.83	67CE1	25030 670н	30.56
67CE1	25030 67CG	30.00	67CE1	25030 67CE2	29.39
660	25030 67CE	68.13	660	25030 67CZ	86.99
660	25030 66C	13.74	660	25030 67CG	55.33
660	25030 67CE	2 73.81	67CD1	25030 67CZ	32.19
67CD1	25030 66C	61.85	67CD1	25030 670н	47.68

			T	ABLE XV			
67CD1	25030	67CG	16.18	67CD1	25030	67CE2	34.38
67CZ	25030	66C	75.55	67CZ	25030	670H	16,91
67CZ	25030	67CG	34.81	67CZ	25030	67CE2	
66C	25030	670H	90.11	66C	25030	67CG	47.06
66C	25030	67CE2	61.38	670H	25030	67CG	51.72
670H	25030	67CE2	28.75	67CG	25030	67CE2	28.79
1610	25C31	161C	15.89	1610	25C31	163CB	92.78
1610	25C31	163N	61.31	1610	25C31	25 <i>S</i> G	64.02
1610	25C31	1600	63.87	1610	25C31	161CA	29.15
1610	25C31	162N	24.80	1610	25C31	162C	50.12
1610	25C31	162CA	32.14	161C	25C31	163CB	90.31
161C	25C31	163N	58.98	161C	25C31	25 <b>S</b> G	74.60
161C	25C31	1600	51.49	161C	25C31	161CA	18.45
161C	25C31	162N	14.50	161C	25C31	162C	44.52
161C	25C31	162CA	29.60	660	25C31	163CB	84.58
660	25C31	25SG	96.91	163CB	25C31	163N	31.73
163CB	25C31	25SG	54.53	163CB	25C31	162N	76.07
163CB	25C31	162C	45.95	163CB	25C31	162CA	61.61
163N	25C31	25SG	46.05	163N	25C31	1600	95.54
163N	25C31	L61CA	75.21	163N	25C31	162N	45.16
163N	25C31	162C	16.28	163N	25C31	162CA	29.89
25 <i>S</i> G	25C31 1	L61CA	92.28	25SG	25C31	162N	68.65
25 <i>S</i> G	25C31	L62C	56.25	25 <i>S</i> G	25C31	162CA	53.18
1600	25C31 1	L61CA	34.72	1600	25C31	162N	58.12
1600	25C31 1	L62C	79.70	1600	25C31	162CA	74.94
161CA	25C31 1	162N	30.24	161CA	25C31	162C	59.56
161CA	25C31 1	62CA	47.15	162N	25C31	162C	30.15
162N	25C31 1	.62CA	17.36	162C	25C31	162CA	18.28
660	25C32	66C	6.90	660	25C32	26CB	47.54
660	25C32	66N	33.71	660	25C32	67CA	32.46
660	25C32	67N	16.73	660	25C32	68SD	78.15
660	25C32	66CA	18.01	66C	25C32	26CB	54.44
66C	25C32	66N	33.07	66C	25C32	67CA	31.49
66C	25C32	67N	13.91	66C	25C32	68SD	82.32
66C	25C32	66CA	15.84	163CB	25C32	26CB	63.43
163CB	25C32 1	610	78.07	1.63CB	25C32	68SD	47.45
25CE	25C32	66N	59.39	26CB	25C32	67 <b>CA</b>	62.90
26CB	25C32	67N	59.03	26CB	25C32	68SD	57.55

			T	ABLE XV			
26CB	25C32	2 66CA	57.10	66N	25C32	67CA	64.54
66N	25C32	2 67N	46.76	66N	25C32	66CA	17.45
67CA	25C32	2 67N	17.89	67CA	25C32	68SD	59.36
67CA	25C32	66CA	47.17	67N	25C32	68SD	73.47
67N	25C32	66CA	29.33	68SD	25C32	66CA	96.00
660	25C33	68SD	88.15	660	25C33	67CA	38.20
660	25C33	66C	8.09	660	25C33	67CD1	67.84
660	25C33	26CB	42.16	68SD	25C33	163CB	58.04
68SD	25C33	209CD2	90.24	68SD	25C33	68CE	24.67
68SD	25C33	134CB	82.77	68SD	25C33	67CA	68.81
68SD	25C33	66C	87.57	68SD	25C33	67CD1	97.64
68SD	25C33	163CA	69.15	68SD	25C33	26CB	61.04
68SD	25C33	163N	85.62	163CB	25C33	68CE	66.68
163CB	25C33	134CB	76.70	163CB	25C33	163CA	16.69
163CB	25C33	26CB	62.05	163CB	25C33	163N	28.80
209CD2	25C33	68CE	67.71	209CD2	25C33	134CB	48.90
209CD2	25C33	67CA	91.59	209CD2	25C33	67CD1	58.37
68CE	25C33	134CB	60.70	68CE	25C33	67CA	80.55
68CE	25C33	67CD1	93.40	68CE	25C33	163CA	71.10
68CE	25C33	26CB	85.68	68CE	25C33	163N	88.13
134CB	25C33	163CA	63.97	134CB	25C33	163N	69.60
67CA	25C33	66C	31.31	67CA	25C33	67CD1	43.75
67CA	25C33	26CB	61.06	6 <b>6</b> C	25C33	67CD1	59.98
66C	25C33	26CB	47.92	163CA	25C33	26CB	77.85
163CA	25C33	163N	17.21	26CB	25C33	163N	81.08
134CB	25C34	163CB	98.21	134CB	25C34	209CD2	58.87
134CB	25C34	134CA	20.10	134CB	25C34	163CA	84.05
134CB	25C34	163N	93.22	134CB	25C34	68SD	90.66
134CB	25C34	1620	68.42	134CB	25C34	162C	84.53
134CB	25C34	68CE	68.29	134CB	25C34	1600	90.59
134CB	25C34	1330	48.20	134CB	25C34	134C	23.47
134CB	25C34	134N	22.08	163CB	25C34	134CA	80.28
163CB	25C34	163CA	22.29	2630B	25C34	163N	35.89
163CB	25C34	68SD	54.05	163CB	25C34	1620	61.05
163CB	25C34	162C	53.04	163CB	25C34	68CE	66.68
163CB	25C34	1610	82.47	163CB	25C34	660	82.69
163CB	25C34	161C	97.58	163CB	25034	1330	51.04
163CB	25C34	134C	88.38	163CB	25C34	134N	76.48

•			Τ.	ABLE XV			
209CD2	25C34	134CA	78.36	209CD2	25C34	68SD	85.07
209CD2	25C34	68CE	68.09	209CD2		1600	89.46
209CD2	25C34	660	98.98	209CD2		1330	91.98
209CD2	25C34	134C	80.67	209CD2	25C34		73.48
134CA	25C34	163CA	64.27	134CA		163N	73.36
134CA	25C34	68SD	87.47	134CA	25C34		50.99
134CA	25C34	162C	66.34	134CA	25C34	68CE	68.94
134CA	25C34	1600	96.11	134CA		161C	93.44
134CA	25C34	1330	34.17	134CA	25C34		14.63
134CA	25C34	134N	13.38	163CA	25C34		20.60
163CA	25C34	68SD	71.18	163CA	25C34	1620	39.10
163CA	25C34	162C	33.94	163CA	25C34	68CE	76.73 <sup>.</sup>
163CA	25C34	1610	73.90	163CA	25C34	161C	74.17
163CA	25C34	1330	43.83	163CA	25C34	134C	69.19
163CA	25C34	134N	64.70	163N	25C34	68SD	89.44
163N	25C34	1620	30.77	163N	25C34	162C	17.88
163N	25C34	68CE	97.26	163N	25C34	1600	97.64
163N	25C34	1610	53.38	163N	25C34	161C	53.96
163N	25C34	1330	61.87	163N	25C34	134C	73.07
163N	25C34	134N	77.98	68SD	25C34	68CE	23.41
68SD	25C34	660	65.12	68SD	25C34	L330	57.26
68SD	25C34	134N	74.79	1620	25C34 1	L62C	16.36
1620	25C34	1600	80.07	1620	25C34	1610	57.49
1620		161C	48.98	1620	25C34 1	1330	57.78
1620	25C34	134C	45.59	1620	25C34	L34N	60.64
162C		1600	82.14	162C	25C34 1	L <b>61</b> 0	46.60
162C		161C	42.17	162C	25C34 1	1330	66.02
162C	25C34		61.91	162C		.34N	74.59
68CE	25C34		84.63	68CE	25C34 1		46.00
68CE	25C34		83.30	68CE	25C34 1		55.59
1600	25C34		52.91	1600	25C34 1	.61C	45.36
1600	25C34		81.68	1610	25C34		93.09
1610	25C34		14.42	1610	25C34 1		94.94
161C	25C34		81.60	1330	25C34 1		47.42
1330	25C34		26.12	134C	25C34 1		27.76
209CD2	25C35	67CD1	84.45	209CD2	25C35	67CE1	80.21
209CD2	25C35	68SD	37.07	209CD2	25C35 2		14.43
209CD2	25 <b>C</b> 35	68CE	73.57	209CD2	25C35 2	340H2	69.51

			TA	BLE XV			
209CD2	25C35 1	.34CB	48.83	209CD2	25C35	67CZ	91.21
67CD1	25C35	660	85.63	67CD1	25C35		57.17
67CD1	25C35	67CE1	19.94	67CD1	25C35		19.64
67CD1	25C35	67CB	37.21	67CD1	25C35		80.87
67CD1	25C35	66C	72.49	67CD1	25C35	<del>-</del>	56.52
67CD1	25C35	67N	60.31	67CD1	25C35		77.14
67CD1	25C35	67C	68.08	67CD1	25C35		24.12
67CD1	25C35	67CD2	23.57	660	25C35		43.70
660	25C35	67CE1	93.34	660	25C35		67.30
660	25C35	67CB	61.47	660	25C35		79.58
660	25C35	66C	13.35	660	25C35		97.49
660	25C35	67N	29.17	660	25C35		61.59
660	25C35	67C	49.59	660	25C35	67CZ	83.31
660	25C35	57CD2	62.47	67CA	25C35		74.54
67CA	25C35	57CG	38.46	67CA	25C35	67CB	21.40
67CA	25C35	8SD	71.28	67CA	25C35	68CE	86.07
67CA	25C35	6C	33.51	67CA	25C35	2340н2	54.58
67CA	25C35 6	57 <b>N</b>	17.55	67CA	25C35	6 <b>8N</b>	31.22
67CA	25C35 6	57C	15.46	67CA	25C35	67CZ	70.81
67CA	25C35 6	7CD2	42.99	67CE1	25C35	67CG	36.15
67CE1	25C35 6	7CB	56.18	67CE1	25C35	209CG	81.63
67CE1	25C35 6	6C	81.58	67CE1	25C35	2340H2	73.50
67CE1	25C35 6	7 <b>N</b>	73.49	67CE1	25C35	68N	96.98
67CE1	25C35 6	7C	86.96	67CE1	25C35	67CZ	11.55
67CE1	25C35 6	7CD2	33.63	67CG	25C35	67CB	21.11
67CG	25C35 20	9CG	95.25	67CG	25C35	66C	53.96
67CG	25C35 23	40H2	55.92	67CG	25C35	67N	40.77
67CG		8N	62.93	67CG	25C35	67C	51.10
67CG	25C35 6	7CZ	34.20	67CG	25C35	67CD2	11.75
67CB		8SD	84.13	67CB	25C35	209CG	94.85
67CB	25C35 6	8CE	92.04	67CB	25C35	66C	49.06
67CB	25C35 23	40H2	43.66	67CB	25C35	67N	32.31
67CB	25C35 6	8N	41.83	67CB	25C35	67C	30.89
67CB	25C35 6	7CZ	55.30	67CB	25C35	67CD2	30.26
6 <b>8S</b> D	25C35 20	9CG	86.31	68SD	25C35	68CE	23.56
68SD	25C35 6	6C	84.14	68 <b>S</b> D	25C35	2340H2	65.58
68SD	25C35 6	7 <b>N</b>	81.24	68SD	25C35	134CB	74.35
68SD	25C35 68	BN	42.38	68SD	25 <b>C</b> 35	67C	56.14

			Т	ABLE XV			
209CG	25C35	68CE	63.48			2340H2	55.92
209CG	25C35	134CB	53.71	209CG	25C35	68N	92.87
209CG	25C35	67CZ	93.16	68CE	25C35	2340H2	58.89
68CE	25C35	67N	99.75	68CE	25C35		57.04
68CE	25C35	68N	54.84	68CE	25C35		70.76
66C	25C35	2340H2	88.09	66C	25C35		16.98
66C	25C35	68N	57.38	66C	25C35	67C	42.97
66C	25C35	67CZ	72.32	66C	25C35		49.60
2340H2	25C35	67N	71.60	2340H2	25C35	134CB	98.59
2340H2	25C35	68 <b>N</b>	41.88	2340H2	25C35	67C	48.67
2340H2	25C35	67CZ	80.56	2340H2	25C35	67CD2	67.67
67N	25C35	68N	46.13	67N	25C35	67C	30.25
67N	25C35	67CZ	66.52	67N	25C35	67CD2	39.87
68N	25C35	67C	16.06	68N	25C35	67CZ	97.13
68N	25C35	67CD2	71.07	67C	25C35	67CZ	84.80
67C	25C35	67CD2	57.44	67CZ	25C35	67CD2	27.90
1610	25C36	25 <i>S</i> G	73.45	1610	25C36	161C	9.23
1610	25C36	163N	54.83	25SG	25C36	161C	78.40
25SG	25C36	65CA	99.90	25 <i>S</i> G	25C36	26CD1	71.50
25 <i>S</i> G	25C36	163N	45.03	25SG	25C36	26CB	68.53
660	25C36	66N	37.32	660	25C36	65CA	68.43
660	25C36	26CD1	49.39	660	25C36	26CB	39.54
161C	25C36	163N	52.99	66N	25C36	65CA	31.50
6 <b>6N</b>	25C36	26CD1	39.27	6 <b>6N</b>	25C36	26CB	55.35
65CA	25C36	26CD1	45.88	65CA	25C36	26CB	73.66
26CD1	25C36	26CB	29.63	163N	25C36	26CB	78.56
66N	25037	65CA	42.38	66N	25037	660	46.15
66N	25037	65C	19.40	66N	25037	66CA	14.20
66N	25037	66C	33.73	6 <b>6N</b>	25037	26CD1	46.03
66N	25037	65N	50.68	6 <b>6N</b>	25037	640	71.48
66N	25037	26CB	60.88	66N	25037	26CG	47.75
65CA	25037	660	87.59	65CA	25037	65C	23.08
65CA	25037	66CA	56.55	65CA	25037	66C	75.97
65CA	25037	26CD1	55.04	65CA	25037	65N	8.31
65CA	25037	640	33.61	65CA	25037	26CB	84.88
65CA	25037	26CG	67.58	660	25037	65C	65.32
660	25037	66CA	32.98	660	25037	66C	13.52
660	25037	26CD1	56.53	660	25037	65N	95.80

	•	Т	ABLE XV		
660	25037 26CB	40.45	660	25037 26CG	44.04
65C	25037 66CA	33.51	65C	25037 66C	53.13
65C	25037 26CD1	48.33	65C	25037 65N	31.35
65C	25037 640	52.95	65C	25037 26CB	72.24
65C	25037 26CG	56.09	1610	25037 25SG	57.00
66CA	25037 66C	19.92	66CA	25037 26CD1	
66CA	25037 65N	64.84	66CA	25037 640	84.47
66CA	25037 26CB	56.17	66CA	25037 26CG	47.05
66C	25037 26CD1	55.02	66C	25037 65N	84.27
66C	25037 26CB	47.54	66C	25037 26CG	45.65
26CD1	25037 25SG	70.87	26CD1	25037 65N	59.66
26CD1	25037 640	86.49	26CD1	25037 26CB	31.10
26CD1	25037 26CG	15.01	25 <i>S</i> G	25037 26CB	64.40
25 <i>S</i> G	25037 26CG	71.37	65 <b>N</b>	25037 640	27.18
65N	25037 26CB	90.35	65N	25037 26CG	73.15
26CB	25037 26CG	17.30	1610	25N38 161C	6.07
1610	25N38 162CA	37.32	1610	25N38 163N	66.42
1610	25N38 162N	19.53	1610	25N38 162C	50.16
1610	25N38 161CA	13.58	1610	25N38 163CB	91.46
25SG	25N38 162CA	69.08	25 <b>S</b> G	25N38 163N	53.99
25 <i>S</i> G	25N38 162N	87.78	25 <b>S</b> G	25N38 25CB	2.51
25 <i>S</i> G	25N38 162C	65.56	25SG	25N38 163CB	56.74
161C	25N38 162CA	34.04	161C	25N38 163N	61.45
161C	25N38 162N	15.39	161C	25N38 162C	45.21
161C	25N38 161CA	14.75	161C	25N38 163CB	85.67
162CA	25N38 163N	32.71	162CA	25N38 162N	18.81
162CA	25N38 25CB	71.19	162CA	25N38 162C	18.79
162CA	25N38 161CA	48.73	162CA	25N38 163CB	62.05
163N	25N38 162N	47.03	163N	25N38 25CB	56.49
163N	25N38 162C	16.27	163N	25N38 161CA	74.28
163N	25N38 163CB	29.78	162N	25N38 25CB	89.84
162N	25N38 162C	30.82	162N	25N38 161CA	29.96
162N	25N38 163CB	73.48	25CB	25N38 162C	68.00
25CB	25N38 163CB	58.75	162C	25N38 161CA	58.33
162C	25N38 163CB	43.86	161CA	25N38 163CB	95.28
25 <i>S</i> G	25N39 1610	88.31	25 <b>SG</b>	25N39 230	88.64
25 <i>S</i> G	25N39 25CB	14.83	25 <b>S</b> G	25 <b>N39 23C</b>	82.58
25SG	25N39 161C	85.92	25 <i>\$</i> G	25N39 25N	41.66

		т	ABLE XV		
25 <i>S</i> G	25N39 23CA	91.29	25SG	25N39 25CA	24.80
25 <i>S</i> G	25N39 162CA	58.01	1610	25N39 25CB	99.78
1610	25N39 161C	3.40	1610	25N39 162CA	31.86
230	25N39 25CB	77.70	230	25N39 23C	14.66
230	25N39 65CA	50.70	230	25N39 25N	47.04
230	25N39 23CA	29.83	230	25N39 25CA	63.85
25CB	25N39 23C	69.66	25CB	25N39 161C	97.02
25CB	25N39 25N	31.35	25CB	25N39 23CA	76.81
25CB	25N39 25CA	16.71	25CB	25N39 162CA	68.20
23C	25N39 65CA	64.77	23C	25N39 25N	41.89
23C	25N39 23CA	18.46	23C	25N39 25CA	58.51
161C	25N39 162CA	28.91	65CA	25N39 25N	88.24
65CA	25N39 23CA	73.18	25N	25N39 23CA	54.84
25N	25N39 25CA	16.92	25N	25N39 162CA	99.14
23CA	25N39 25CA	69.81	25CA	25N39 162CA	82.61
1600	25N40 1610	83.18	1600	25N40 161C	65.67
1600	25N40 161CA	44.84	1600	25N40 160C	14.22
1600	25N40 161N	30.23	1600	25N40 162N	67.90
1610	25N40 161C	19.36	1610	25N40 161CA	38.54
1610	25N40 160C	72.40	1610	25N40 161N	54.55
1610	25N40 162N	26.72	161C	25N40 161CA	23.35
161C	25N40 160C	53.79	161C	25N40 161N	35.96
161C	25N40 162N .	13.14	161CA	25N40 160C	35.76
161CA	25N40 161N	19.52	161CA	25N40 162N	31.98
160C	25N40 161N	17.87	160C	25N40 162N	54.39
161N	25N40 162N	37.86			

**TABLE XVI** 

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
1840	25C1	184CD1	70.91	1840	25C1	184CG	59.71
1840	25Ci	184CB	40.07	1840	25C1	184CA	32.41
1840	25C1	180D1	51.11	1840	25C1	184C	14.14
1840	25C1	184NE1	85.23	1840	25C1	184CD2	70.75
184CD1	25C1	184CG	18.26	184CD1	25C1	184CB	34.36
184CD1	25C1	184CA	39.63	184CD1	25C1	180D1	88.02
184CD1	25C1	184C	58.65	184CD1	25C1	184NE1	15.36
184CD1	25C1	184CD2	26.43	184CG	25C1	184CB	19.78
184CG	25C1	184CA	33.68	184CG	25C1	180D1	90.72
184CG	25C1	184C	50.19	184CG	25C1	184NE1	27.27
184CG	25C1	184CD2	15.55	184CB	25C1	184CA	19.99
184CB	25C1	180D1	77.76	184CB	25C1	184C	31.92
184CB	25C1	184NE1	46.53	184CB	25C1	184CD2	31.52
184CA	25C1	180D1	58.36	184CA	25C1	184C	19.14
184CA	25C1	184NE1	54.76	184CA	25C1	184CD2	48.54
180D1	25C1	184C	48.87	184C	25C1	184NE1	73.60
184C	25C1	184CD2	63.42	184NE1	25C1	184CD2	26.12
200	25C2	20C	16.36	200	25C2	21NE2	67.96
200	25C2	20N	38.80	200	25C2	20CA	32.93
200	25C2	19CG	52.37	200	25C2	180D1	81.91
20C	25C2	21NE2	65.02	20C	25C2	20 <b>N</b>	33.58
20C	25C2	20CA	20.23	20C	25C2	19CG	62.24
20C	25C2	180D1	73.06	21NE2	25C2	20 <b>N</b>	97.72
21NE2	25C2	20CA	79.83	184CD1	25C2	20N	95.12
184CD1	25C2	19CG	55.67	184CD1	25C2	180D1	86.02
184CD1	25C2	184NE1	16.52	184CD1	25C2	184CG	16.21
184CD1	25C2	184CA	36.11	20N	25C2	20CA	19.09
20N	25C2	19CG	44.54	20N	2 F C 2	180D1	43.33
20N	25C2	184CA	81.17	20CA	25C2	19CG	60.59
20CA	25C2	180D1	53.05	20CA	25C2	184CA	98.88

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-			TA	BLE XVI			
19CG	25C2	180D1	66.67	19CG	25C2	184NE1	60.03
19CG	25C2	184CG	69.53	19CG	25C2	184CA	62.53
180D1	25C2	184CG	83.94	180D1	25C2	184CA	53.91
184NE1	25C2	184CG	27.27	184NE1	25C2	184CA	52.47
184CG	25C2	184CA	30.07	200	25C3	21NE2	68.98
200	25C3	20C	12.74	200	25C3	19CG	52.30
200	25C3	19CD	62.76	184CD1	25C3	184NE1	19.43
184CD1	25C3	19CG	57.07	184CD1	25C3	184CG	15.46
184CD1	25C3	184CE2	27.70	184CD1	25C3	19CD	51.26
184NE1	25C3	19CG	64.69	184NE1	25C3	184CG	28.26
184NE1	25C3	184CE2	15.28	184NE1	25C3	19CD	52.93
21NE2	25C3	20C	62.81	20C	25C3	19CG	59.85
20C	25C3	19CD	72.69	19CG	25C3	184CG	69.91
19CG	25C3	184CE2	79.68	19CG	25C3	19CD	17.58
184CG	25C3	184CE2	27.82	184CG	25C3	19CD	66.23
184CE2	25C3	19CD	68.19	184NE1	25C4	184CD1	21.40
184NE1	25C4	184CE2	20.66	184NE1	25C4	184CG	31.78
184NE1	25C4	184CD2	31.15	184NE1	25C4	184CZ2	34.27
184CD1	25C4	184CE2	34.03	184CD1	25C4	184CG	18.28
184CD1	25C4	184CD2	31.27	184CD1	25C4	184CZ2	51.25
184CE2	25C4	184CG	32.35	184CE2	25C4	184CD2	18.86
184CE2	25C4	184CZ2	17.60	184CG	25C4	184CD2	19.32
184CG	25C4	184CZ2	49.25	184CD2	25C4	184CZ2	32.53
184CE2	25C5	184NE1	21.44	184CE2	25C5	184CD2	21.99
184CE2	25C5	184CD1	35.25	184CE2	25C5	1.84CG	36.03
184CE2	25C5	184CZ2	17.88	184CE2	25C5	184CE3	33.85
184CE2	25C5	184CB	53.93	184CE2	25C5	184CH2	28.11
184CE2	25C5	184CZ3	34.04	184NE1	25C5	184CD2	35.23
184NE1	25C5	184CD1	21.39	184NE1	25C5	184CG	35.13
184NE1	25C5	184CZ2	35.43	184NE1	25C5	184CE3	51.78
184NE1	25C5	184CB	52.94	184NE1	25C5	184CH2	48.87
184NE1	25C5	184CZ3	55.18	184CD2	25C5	184CD1	35.19
184CD2	25C5	184CG	22.35	184CD2	25C5	184CZ2	34.58
184CD2	25C5	184CE3	17.91	184CD2	25C5	184CB	36.19
184CD2	25C5	184CH2	34.79	184CD2	25C5	184CZ3	
184CD1	25C5	184CG	21.04	184CD1	25C5	184CZ2	52.70
184CD1	25C5	184CE3	52.87	184CD1	25C5	184CB	34.98
134CD1	25C5	184CH2	62.57	184CD1	25C5	184CZ3	62.67

			TAI	BLE XVI			
184CG	25C5	184CZ2	53.00	184CG	25C5	184CE3	36.87
184CG	25C5	184CB	18.54	184CG	25C5	184CH2	56.68
184CG	25C5	184CZ3	50.46	184CZ2	25C5	184CE3	38.17
184CZ2	25C5	184CB	69.96	184CZ2	25C5	184CH2	15.84
184CZ2	25C5	184CZ3	29.77	184CE3	25C5	184CB	44.40
184CE3	25C5	184CH2	29.96	184CE3	25 <b>C</b> 5	184CZ3	16.03
184CB	25C5	184CH2	70.55	184CB	25C5	184CZ3	60.21
184CH2	25C5	184CZ3	16.60	184CG	25C6	184CD1	20.02
184CG	25C6	184CB	21.95	184CG	25C6	184CD2	20.42
184CG	25C6	1840	61.94	184CG	25C6	184NE1	30.53
184CG	25C6	184CE2	30.58	184CG	25 <b>C</b> 6	184CA	33.96
184CG	25C6	184CE3	33.66	184CG	25C6	184C	50.16
184CD1	25C6	184CB	37.32	184CD1	25C6	184CD2	32.14
184CD1	25C6	1840	71.19	184CD1	25C6	184NE1	17.93
184CD1	25C6	184CE2	29.87	184CD1	25C6	184CA	39.85
184CD1	25C6	184CE3	48.53	184CD1	25C6	184C	57.95
184CB	25C6	184CD2	37.74	184CB	25C6	1840	40.76
184CB	25C6	184NE1	51.79	184CB	25C6	184CE2	51.93
184CB	25C6	184CA	19.22	184CB	25C6	184CE3	43.86
184CB	25C6	184C	30.42	184CD2	25C6	1840	78.21
184CD2	25C6	184NE1	30.21	184CD2	25C6	184CE2	18.12
184CD2	25C6	184CA	53.55	184CD2	25C6	184CE3	16.59
184CD2	25C6	184C	68.06	1840	25C6	184NE1	88.69
1840	25C6	184CE2	92.47	1840	25C6	184CA	31.36
1840	25C6	184CE3	80.46	1840	25C6	184C	13.35
184NE1	25C6	184CE2	17.71	184NE1	25C6	184CA	57.51
184NE1	25C6	184CE3	45.78	184NE1	25C6	184C	75.58
184CE2	25C6	184CA	63.83	184CE2	25C6	184CE3	30.35
184CE2	25C6	184C	80.70	184CA	25C6	184CE3	62.43
184CA	25C6	184C	18.09	184CE3	25C6	184C	73.23
184NE1	25C7	184CE2	21.08	184NE1	25C7	184CZ2	38.55
184NE1	25C7	184CD1	16.84	184NE1	25C7		27.78
184NE1	25C7	184CH2	48.97	184NE1	25C7	184CG	
184CE2	25C7	184CZ2	20.61	184CE2	25C7	184CD1	
184CE2		184CD2	15.35	184CE2	25 <b>C</b> 7	184CH2	
184CE2			26.62	184CZ2	25C7	184CD1	51.38
184CZ2			31.85	184CZ2	25C7	184CH2	
184CZ2			46.61	184CD1	25C7	184CD2	28.53
· <del>-</del> -							

			TA	BLE XVI			
184CD1	25C7	184CH2	59.02	184CD1	25C7	184CG	15.12
184CD2	25C7	184CH2	33.93	184CD2	25C7	184CG	16.91
184CH2	25C7	184CG	50.64	184NE1	2508	184CE2	19.75
184NE1	2508	184CZ2	37.94	184NE1	2508	184CD1	15.31
184NE1	2508	19NE2	71.47	184NE1	2508	19CD	57.15
184CE2	2508	184CZ2	19.95	184CE2	2508	184CD1	31.02
184CE2	2508	19NE2	90.51	184CE2	2508	19CD	76.72
184CZ2	2508	184CD1	50.82	184CZ2	2508	19CD	92.06
184CD1	2508	19NE2	67.22	184CD1	2508	19CD	51.55
19NE2	2508	19CD	15.86	184NE1	25C9	184CZ2	42.18
184NE1	25C9	184CE2	21.04	184NE1	25C9	162ND1	65.62
184NE1	25C9	162CE1	48.16	184NE1	25C9	184CD1	10.29
184NE1	25C9	19NE2	73.80	184NE1	25C9	190E1	48.69
184NE1	25C9	19CD	57.90	184NE1	25C9	184CH2	49.13
184CZ2	25C9	184CE2	21.63	184CZ2	25C9	162ND1	65.18
184CZ2	25C9	162CE1	57.30	184CZ2	25 <b>C</b> 9	184CD1	51.49
184CZ2	25 <b>C</b> 9	190E1	86.41	184CZ2	25C9	19CD	98.49
184CZ2	25C9	184CH2	7.78	184CE2	25C9	162ND1	66.81
184CE2	25 <b>C</b> 9	162CE1	52.84	184CE2	25C9	184CD1	29.90
184CE2	25C9	19NE2	94.57	184CE2	25C9	190E1	68.42
184CE2	25C9	19CD	78.74	184CE2	25C9	184CH2	28.19
162ND1	25C9	162CE1	18.04	162ND1	25 <b>C</b> 9	184CD1	72.19
162ND1	25C9	19NE2	76.32	162ND1	25 <b>C9</b>	190E1	58.44
162ND1	25C9	19CD	71.67	162ND1	25 <b>C</b> 9	184CH2	70.09
162CE1	25C9	184CD1	54.22	162CE1	25C9	19NE2	69.37
162CE1	25C9	190 <b>E</b> 1	46.03	162CE1	25 <b>C9</b>	19CD	60.55
162CE1	25C9	184CH2	63.87	184CD1	25C9	19NE2	66.83
184CD1	25C9	190E1	44.17	184CD1	25 <b>C9</b>	19CD	50.99
184CD1	25C9	184CH2	58.03	19NE2	25 <b>C9</b>	190E1	27.69
19NE2	25C9	19CD	15.92	190E1	25C9	19CD	14.75
190E1	25C9	184CH2	94.16	184NE1	25010	162ND1	90.68
184NE1	25010	162CE1	67.15	184NE1	25010	184CE2	23.02
184NE1	25010	184CZ2	46.79	184NE1	25010	190E1	64.53
184NE1	25010	184CD1	12.02	184NE1	25010	19NE2	90.63
184NE1	25010	19CD	72.30	184NE1	25010	162CG	91.69
184NE1	25010	162NE2	63.41	184NE1	25010	184CD2	16.34
184NE1	25010	184CH2	50.63	184NE1	25010	162CD2	76.58
184NE1	25010	1.84CG	0.86	162ND1	25010	162CE1	25.39

		T	ABLE XVI			
162ND1	25010 18	4CE2 85.65	162ND1	25010	184CZ2	80.71
162ND1	25010 1	90E1 79.25	162ND1	25010	184CD1	93.79
162ND1	25010 1	9NE2 99.71	162ND1	25010	19CD	94.77
162ND1	25010 16	2CG 11.75	162ND1	25010	162NE2	27.28
162ND1	25010 18	4CD2 87.44	162ND1	25010	184CH2	80.29
162ND1	25010 16	2CD2 17.27	162ND1	25010	162CB	25.40
162ND1	25010 2	5CB 48.85	162ND1	25010	184CG	91.42
162ND1	25010 2	5SG 48.57	162CE1	25010	184CE2	67.89
162CE1	25010 18	4CZ2 72.02	162CE1	25010	190E1	61.36
162CE1	25010 18	4CD1 68.78	162CE1	25010	19NE2	90.18
162CE1	25010 1	9CD 78.77	162CE1	25010	162CG	32.27
162CE1	25010 16	2NE2 9.58	162CE1	25010	184CD2	67.64
162CE1	25010 18	4CH2 73.24	162CE1	25010	162CD2	21.07
162CE1	25010 16	2CB 48.98	162CE1	25010	25CB	50.48
162CE1	25010 18	4CG 67.80	162CE1	25010	25SG	61.27
184CE2	25010 18	4CZ2 23.77	184CE2	25010	190E1	85.64
184CE2	25010 18	4CD1 35.04	184CE2	25010	19CD	95.11
184CE2	25010 16	2CG 82.23	184CE2	25010	162NE2	60.72
184CE2	25010 18	4CD2 6.70	184CE2	25010	184CH2	27.61
184CE2	25010 16	2CD2 68.81	184CE2	25010	162CB	92.97
184CE2	25010 18	4CG 23.63	184CZ2	25010	184CD1	58.81
184CZ2	25010 16	2CG 73.16	184CZ2	25010	162NE2	62.79
184CZ2	25010 18	4CD2 30.47		25010	184CH2	3.84
184CZ2	25010 16	2CD2 63.78	3 184CZ2	25010	162CB	78.57
184CZ2	25010 18	4CG 47.41	1.90E1	25010	184CD1	54.02
190E1	25010 1	.9NE2 33.63	5 190E1	25010	19CD	17.62
190E1	25010 16	2CG 90.23	3 190E1	25010	162NE2	68.58
190E1	25010 18	4CD2 79.63	2 190E1	25010	162CD2	82.33
190E1	25010 2	SCB 46.49	9 190E1	25010	184CG	64.24
190El	25010 2	55G 67.15	5 184CD1	25010	19NE2	78.67
184CD1	25010 1	.9CD 60.49	9 184CD1	25010		97.06
184CD1	25010 16	2NE2 67.0	8 184CD1	25010	184CD2	28.35
184CD1	25010 18	4CH2 62.6	4 184CD1	25010		81.91
184CD1	25010 2	SCB 94.2	9 184CD1		184CG	11.41
19NE2	25010 1	.9CD 18.9	7 19NE2		162NE2	98.89
19NE2	25010 2	25CB 52.3	1 19NE2		184CG	90.06
19NE2	25010 2	25SG 64.0	8 19CD	25010		36.19
19CD	25010 18	34CD2 88.5	3 19CD	25010	162CD2	99.61

			TAI	BLE XVI			
19CD	25010	25CB	54.29	19CD	25010	184CG	71.79
19CD	25010	25 <b>S</b> G	71.77	162CG	25010	162NE2	30.49
162CG	25010	184CD2	85.26	162CG	25010	184CH2	72.15
162CG	25010	162CD2	15.20	162CG	25010	162CB	16.86
162CG	25010	25CB	60.05	162CG	25010	184CG	92.50
162CG	25010	25 <b>S</b> G	56.96	162NE2	25010	184CD2	61.36
162NE2	25010	184CH2	63.86	162NE2	25010	162CD2	16.25
162NE2	25010	162CB	47.22	162NE2	25010	25CB	59.84
162NE2	25010	184CG	64.14	162NE2	25010	25 <b>S</b> G	69.18
184CD2	25010	184CH2	34.30	184CD2	25010	162CD2	71.14
184CD2	25010	162CB	97.22	184CD2	25010	184CG	16,94
184CH2	25010	162CD2	63.64	184CH2	25010	162CB	76.54
184CH2	25010	184CG	51.24	162CD2	25010	162CB	31.36
162CD2	25010	25CB	62.62	162CD2	25010	184CG	77.38
162CD2	25010	25SG	65.53	162CB	25010	25CB	66.47
162CB	25010	25 <b>SG</b>	56.57	25CB	25010	25 <b>S</b> G	21.55
1610	25C11	162ND1	77.52	1610	25C11	162CB	48.77
1610	25C11	162CG	66.44	1610	25C11		9.16
1610	25C11	1610D1	44.79	1610	25C11	162CE1	89.91
1610	25C11	162CA	34.68	162ND1	25C11	162CB	36.88
162ND1	25C11	162CG	18.06	162ND1	25C11	184CZ2	57.93
162ND1	25C11	161C	73.11	162ND1	25C11	1610D1	88.11
162ND1	25C11	162CE1	12.39	162ND1	25C11	162CA	43.39
162CB	25C11	162CG	20.08	162CB	25C11	184CZ2	75.16
162CB	25C11	161C	41.40	162CB	25C11	1610D1	51.91
162CB	25C11	162CE1	47.69	162CB		162CA	18.23
162CG	25C11	184CZ2	60.65	162CG	25C11		60.17
162CG	25C11	1610D1	70.16	162CG		162CE1	27.75
162CG		162CA	32.09	184CZ2		1610D1	96.89
			48.52				
			38.12				
			29.74				
			54.92				
1610	25C12	1610D1	53.96				
		161C		1610			
			38.06				
			14.06				
1610D1	25C12	162CB	55.80	161001	25C12	161CB	30.86

			TA	BLE XVI			
1610D1	25C12	162ND1	86.96	161CG	25C12	161C	40.85
161CG	25C12	162CB	65.21	161CG	25C12	161CB	19.00
161CG	25C12	162ND1	96.52	161C	25C12	162CB	40.99
161C	25C12	161CB	31.65	161C	25C12	162ND1	65.80
162CB	25C12	161CB	67.55	162CB	25C12	162ND1	31.38
161CB	25C12	162ND1	96.34	1610D1	25C13	1370	81.25
1610D1	25C13	137C	63.73	1610D1	25C13	138N	56.48
1610D1	25C13	138CA	69.03	1610D1	25C13	137CB	63.42
1610D1	25C13	161CG	11.19	1610D1	25C13	1610	44.97
1610D1	25C13	162CB	53.52	1610D1	25C13	137CA	54.88
1370	25C13	137C	17.61	1370	25C13	138N	31.56
1370	25C13	184CZ2	73.45	1370	25C13	138CA	38.68
1370	25C13	137CB	38.26	1370	25C13	143NE2	53.14
1370	25C13	161CG	88.09	1370	25C13	184CH2	57.10
1370	25C13	162CB	89.51	1370	25C13	137CA	30.26
137C	25C13	138N	18.18	137C	25C13	184CZ2	85.16
137C	25C13	138CA	33.43	137C	25C13	137CB	33.67
137C	25C13	143NE2	68.48	137C	25C13	161CG	70.50
137C	25C13	184CH2	70.29	137C	25C13	162CB	80.53
137C	25C13	137CA	17.92	138N	25C13	138CA	19.51
138N	25C13	137CB	49.22	138N	25C13	143NE2	71.83
138N	25C13	161CG	60.17	138N	25C13	184CH2	87.76
138N	25C13	162CB	88.70	138N	25C13	137CA	30.79
184CZ2	25C13	137CB	62.75	184CZ2	25C13	143NE2	73.88
184CZ2	25C13	184CH2	16.99	184CZ2	25C13	162CB	71.91
184CZ2	25C13	137CA	79.40	138CA	25C13	137CB	66.92
138CA	25C13	143NE2	60.03	139CA	25C13	161CG	69.26
138CA	25C13	184CH2	94.51.	138CA	25C13	137CA	49.37
137CB	25C13	143NE2	87.28	137 <b>CB</b>	25C13	161CG	74.16
137CB	25C13	1610	86.27	137CB		184CH2	53.95
137CB	25C13	162CB	51.39	137 <b>CB</b>	25C13	137CA	18.78
143NE2	25C13	184CH2	61.44	143NE2	25C13	137CA	83.39
161CG	25C13	1610	44.82	161CG	25C13	162CB	62.01
161CG	25C13	137CA	64.10	1610	25C13	162CB	40.52
1610	25C13	137CA	89.27	194CH2	25C13	162CB	78.30
184CH2	25C13	137CA	67.80	162 <b>CB</b>	25C13	137CA	62.63
143NE2	25C14	184CZ2	89.78	143NE2	25014	1370	59.02
143NE2	25C14	184CH2	73.23	143NE2	25C14	143CD	7.66

			TA	BLE XVI			
143NE2	25C14	137C	70.92	143NE2	25C14	138CA	63.43
143NE2	25C14	138N	72.92	184CZ2	25C14	1370	73.56
184CZ2	25C14	184CH2	18.41	184CZ2	25C14	143CD	82.48
184CZ2	25C14	137C	80.26	184CZ2	25C14	138N	95.58
1370	25C14	184CH2	58.32	1370	25C14	143CD	59.13
1370	25C14	137C	14.37	1370	25C14	138CA	35.52
1370	25C14	138N	26.67	184CH2	25C14	143CD	66.37
184CH2	25C14	137C	67.66	184CH2	25C14	138CA	93.75
184CH2	25C14	138N	83.05	143CD	25C14	137C	72.08
143CD	25C14	138CA	68.38	143CD	25C14	138N	76.11
137C	25C14	138CA	29.93	137C	25C14	138N	15.47
138CA	25C14	138N	17.13	1610D1	25C15	138CA	95.49
1610D1	25C15	138N	74.59	1610D1	25C15	137C	77.97
1610D1	25C15	138CB	88.03	1610D1	25C15	1370	95.93
1610D1	25C15	161CG	16.54	1610D1	25C15	137CA	62.55
1610D1	25C15	161ND2	28.48	1610D1	25C15	137N	44.50
1610D1	25C15	137CB	65.90	1610D1	25C15	1380G	75.32
1610D1	25C15	161CB	26.73	1610D1	25C15	1610	41.09
138CA	25C15	138N	26.47	138CA	25C15	137C	42.87
138CA	25C15	138CB	25.16	138CA	25C15	1370	47.79
138CA	25C15	161CG	93.95	138CA	25C15	137CA	58.55
138CA	25C15	161ND2	78.54	138CA	25C15	137N	64.03
138CA	25C15	138C	11.03	138CA	25C15	137CB	75.54
138CA	25C15	143NE2	66.87	138CA	25C15	1380G	29.01
138CA	25C15	1380	24.54	138N	25C15	137C	22.54
138N	25C15	138CB	42.08	138N	25C15	1370	37.92
138N	25C15	161CG	78.54	138N	25C15	137CA	33.52
138N	25C15	161ND2	66.91	138N	25C15	137N	37.71
138N	25C15	138C	32.59	138N	25C15	137CB	52.24
138N	25C15	143NE2	80.31	138N	25C15	1380G	36.11
138N	25C15	161CB	95.16	138N	25C15	1380	46.34
137C	25C15	138CB	63.59	137C	25C15	1370	19.93
137C	25C15	161CG	87.64	137¢	25C15	137CA	18.45
137C	25C15	161ND2	80.75	137C	25C15	137N	33.66
137C	25C15	138C	43.43	137C	25C15	137CB	32.73
137C	25C15	143NE2	72.28	137C	25C15	1.380G	58.64
137C	25C15	1380	53.23	1.38CB	25C15	1370	
138CB	25C15	161CG	80.25	138CB	25C15	137CA	75.11

			TA	BLE XVI			
138CB	25C15	161ND2	63.55	138CB	25C15	137N	73.01
138CB	25C15	138C	33.69	138CB	25C15	137CB	94.23
138CB	25C15	143NE2	84.72	138CB	25C15	1380G	13.06
138CB	25C15	161CB	92.07	138CB	25C15	1380	41.98
1370	25C15	137CA	33.44	1370	25C15	137N	51.60
1370	25C15	138C	43.03	1370	25C15	137CB	37.73
1370	25C15	143NE2	53.68	1370	25C15	1380G	71.49
1370	25C15	1380	46.98	161CG	25C15	137CA	74.83
161CG	25C15	161ND2	16.70	161CG	25C15	137N	56.04
161CG	25C15	137CB	81.22	161CG	25C15	1380G	68.77
161CG	25C15	161CB	16.63	161CG	25C15	1610	45.51
137CA	25C15	161ND2	72.19	137CA	25C15	137N	18.82
137CA	25C15	138C	61.01	137CA	25C15	137CB	19.45
137CA	25C15	143NE2	86.99	137CA	25C15	1380G	66.83
137CA	25C15	161CB	89.10	137CA	25C15	1380	71.60
137CA	25C15	1610	90.57	161ND2	25C15	137N	54.06
161ND2	25C15	138C	89.54	161ND2	25C15	137CB	83.68
161ND2	25C15	1380G	52.25	161ND2	25C15	161CB	30.53
161ND2	25C15	1610	62.17	137N	25C15	138C	69.92
137N	25C15	137CB	31.11	137N	25C15	1380G	61.80
137N	25C15	161CB	70.66	137N	25C15	1380	82.86
137N	25C15	1610	77.59	138C	25C15	137CB	75.69
138C	25C15	143NE2	55.85	138C	25C15	1380G	39.61
138C	25C15	1380	14.34	137CB	25C15	143NE2	85.14
137CB	25C15	1380G	86.28	137CB	25C15	161CB	91.92
137CB	25C15	1380	83.44	137CB	25C15	1610	82.09
143NE2	25C15	1380G	94.34	143NE2	25C15	1380	43.49
1380G	25C15	161CB	82.00	1380G	25C15	1380	50.85
161CB	25C15	1610	34.58	1610	25C16	162ND1	76.53
1610	25C16	161C	3.98	1610	25C16	162CB	45.40
1610	25C16	25 <b>S</b> G	72.69	1610	25C16	162CA	34.24
1610	25C16	162CG	62.41	1610	25C16	162N	17.52
162ND1	25C16	161C	72.86	162ND1	25C16	162CB	33.34
162ND1	25C16	253G	49.81	162ND1	25C16	162CA	43.06
162ND1	25C16	162CG	15.65	162ND1	25C16	162N	59.06
161C	25C16	162CB	41.47	161C	25C16	25SG	71.63
161C	25C15	162CA	30.96	161C	25C16	162CG	58.58
161C	25C16	162N	14.06	162CB	25C16	25SG	60.72

		TA	BLE XVI		
162CB	25C16 162C	A 19.06	162CB	25C16 162CG	17.85
162CB	25C16 162N	29.15	25SG	25C16 162CA	49.45
25SG	25C16 162C	54.90	25SG	25C16 162N	61.48
162CA	25C16 162C	30.90	162CA	25C16 162N	16.94
162CG	25C16 162N	45.31	1610	25017 161C	2.40
1610	25N18 25S	94.42	1610	25N18 162ND1	86.25
1610	25N18 162C	A 41.79	1610	25N18 162CB	49.55
1610	25N18 161C	8.51	1610	25N18 162CG	69.44
1610	25N18 162C	E1 97.59	1610	25N18 162N	24.49
25 <i>S</i> G	25N18 162N	D1 65.37	25SG	25N18 162CA	62.23
25SG	25N18 162C	B 74.73	25SG	25N18 161C	86.60
25SG	25N18 162C	G 67.87	25SG	25N18 25CB	21.50
25 <i>S</i> G	25N18 162C	E1 64.64	25SG	25N18 162N	73.85
25SG	25N18 19N	E2 67.61	25SG	25N18 23CA	79.71
162ND1	25N18 162C	A 50.07	162ND1	25N18 162CB	36.70
162ND1	25N18 161C	79.92	162ND1	25N18 162CG	16.82
162ND1	25N18 25C	B 53.33	162ND1	25N18 162CE1	11.34
162ND1	25N18 162N	65.85	162ND1	25N18 19NE2	73.54
162CA	25N18 162C	B 21.06	162CA	25N18 161C	33.62
162CA	25N18 162C	G 34.82	162CA	25N18 25CB	70.27
162CA	25N18 162C	E1 60.64	162CA	25N18 162N	17.53
162CB	25N18 161C	43.55	162CB	25N18 162CG	19.91
162CB	25N18 25C	B 75.42	162CB	25N18 162CE1	48.04
162CB	25N18 162N	31.25	161C	25N18 162CG	63.15
161C	25N18 162C	E1 91.21	161C	25N18 162N	16.15
162CG	25N18 25C	B 61.89	162CG	25N18 162CE1	28.15
162CG	25N18 162N	49.49	162CG	25N18 19NE2	89.96
25CB	25N18 162C	E1 48.89	25CB	25N18 162N	85.68
25CB	25N18 19N	E2 49.72	25 <b>CB</b>	25N18 23CA	76.32
162CE1	25N18 162N	76.89	162CE1	25N18 19NE2	
19NE2	25N18 23C	A 47.71	25 <b>SG</b>		93.76
25SG	25C19 23C	A 98.71	25 <b>SG</b>		19.06
25 <b>S</b> G	25C19 162N	D1 55.42	25 <b>S</b> G	25C19 23C	83.02
25SG	25C19 162C	A 57.95	25SG	25C19 161C	85.45
25 <b>S</b> G	25C19 23C	84.21	25 <b>SG</b>		39.84
1610	25C19 162N	D1 68.34	1610		
1610	25019 1610	9.82	23CA		
23CA	25019 230	19.53	23CA	25C19 230	30.41

			TA	BLE XVI			
23CA	25C19	25N	58.86	25CB	25C19	162ND1	49.56
25CB	25C19	23C	74.71	25CB	25C19	162CA	67.81
25CB	25C19	161C	97.94	25CB	25C19	230	80.56
25CB	25C19	25N	30.79	162ND1	25C19	162CA	42.46
162ND1	25C19	161C	67.39	162ND1	25C19	25N	78.08
23C	25C19	230	15.10	23C	25C19	25N	44.37
162CA	25C19	161C	30.60	162CA	25C19	25 <b>N</b>	96.99
230	25C19	25N	49.95	184CZ2	25N20	162ND1	57.46
184CZ2	25N20	184NE1	33.66	184CZ2	25N20	184CE2	16.92
184CZ2	25N20	162CE1	48.47	162ND1	25N20	184NE1	53.53
162ND1	25N20	184CE2	56.14	162ND1	25N20	162CE1	14.39
184NE1	25 <b>N</b> 20	184CE2	17.07	184NE1	25N20	162CE1	39.26
184CE2	25N20	162CE1	43.67	1610	25C21	25 <b>S</b> G	97.39
1610	25C21	161C	15.35	1610	25C21	162CA	39.27
1610	25C21	162N	27.81	1610	25C21	161CA	24.14
1610	25C21	25CB	98.96	25 <b>SG</b>	25C21	161C	94.07
25SG	25C21	162CA	60.19	25SG	25C21	162N	78.17
25SG	25C21	65CA	98.56	25SG	25C21	25CB	7.84
161C	25C21	162CA	33.93	161C	25C21	162N	16.70
161C	25C21	161CA	15.71	161C	25C21	25CB	97.60
162CA	25C21	162N	19.15	162CA	25C21	161CA	48.75
162CA	25C21	25CB	63.75	162N	25C21	161CA	30.04
162N	25C21	25CB	82.42	65CA	25C21	25CB	98.99
25 <b>S</b> G	25C22	25CB	34.66	25 <b>SG</b>	25C22	25N	70.64
25SG	25C22	25CA	48.75	25SG	25C22	19NE2	98.64
25SG	25C22	24C	78.81	25 <b>SG</b>	25C22	162ND1	56.90
25SG	25C22	26N	47.65	25 <b>SG</b>	25C22	25C	40.82
25 <b>S</b> G	25C22	1610	81.68	25 <b>SG</b>	25C22	24CA	97.30
25 <b>S</b> G	25C22	162CA	49.19	25 <b>SG</b>	25C22	163N	21.43
25 <b>S</b> G	25C22	26CD1	82.19	25 <b>CB</b>	25C22	25N	45.70
25CB	25C22	25CA	23.52	25CB	25C22	24N	85.13
25CB	25C22	19NE2	64.70	25 <b>CB</b>	25C22	24C	56.87
25CB	25C22	162ND1	56.29	25CB	25C22	26N	47.43
25CB	25C22	25C	31.07	25CB	25C22	24CA	75.01
25CB	25C22	162CA	74.58	25CB	25C22	163N	53.55
25CB	25C22	26CD1	84.62	25N	25C22	25CA	23.10
25N	25C22	23C	59.53	25N	25C22	23CA	76.19
25N	25C22	230	65.52	25 <b>N</b>	25C22	24N	41.74

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			TA	BLE XVI			
25N	25C22	19NE2	55.79	25N	25C22	24C	11.64
25N	25C22	162ND1	98.55	25N	25C22	26N	37.79
25N	25C22	25C	31.73	25N	25C22	24CA	29.34
25N	25C22	163N	92.04	25N	25C22	26CD1	53.86
25CA	25C22	23C	82.62	25CA	25C22	23CA	98.07
25CA	25C22	230	87.79	25CA	25C22	24N	64.51
25CA	25C22	19NE2	62.14	25CA	25C22	24C	33.67
25CA	25C22	162ND1	79.26	25CA	25C22	26N	32.62
25CA	25C22	25C	17.41	25CA	25C22	24CA	52.26
25CA	25C22	162CA	95.54	25CA	25C22	163N	70.10
25CA	25C22	26CD1	64.67	23C	25C22	23CA	22.66
23C	25C22	230	17.69	23C	25C22	24N	19.11
23C	25C22	19NE2	64.93	23C	25C22	24C	49.94
23C	25C22	26N	82.64	23C	25C22	25C	87.50
23C	25C22	24CA	31.15	23C	25C22	26CD1	58.43
23CA	25C22	230	35.06	23CA	25C22	24N	35.02
23CA	25C22	19NE2	59.32	23CA	25C22	24C	68.76
23CA	25C22	24CA	51.14	23CA	25C22	26CD1	80.22
230	25C22	24N	32.37	230	25C22	19NE2	82.31
230	25C22	24C	54.24	230	25C22	26N	77.95
230	25C22	25C	87.44	230	25C22	24CA	36.53
230	25C22	26CD1	46.14	24N	25C22	19NE2	52.35
24N	25C22	24C	33.74	24N	25C22	26N	70.95
24N	25C22	25C	71.93	24N	25C22	24CA	17.11
24N	25C22	26CD1	57.95	19NE2	25C22	24C	60.62
19NE2	25C22	162ND1	74.96	19NE2	25C22	26N	91.86
19NE2	25C22	25C	79.36	19NE2	25C22	24CA	60.80
24C	25C22	26 <b>N</b>	39.04	24C	25C22	25C	38.21
24C	25C22	24CA	18.87	24C	25C22	163N	99.80
24C	25C22	26CD1	45.21	162ND1	25C22	26N	99.77
162ND1	25C22	25C	85.70	162ND1	25C22	1610	61.42
162ND1	25C22	162CA	41.62	162ND1	25022	163N	51.67
26N	25C22	25C	16.78	26 <b>N</b>	25C22	24CA	53.95
26N	25C22	162CA	94.45	26 <b>N</b>	25C22	163N	65.37
26N	25C22	26CD1	38.29	25C	25022	24CA	56.48
25C	25C22	162CA	90.01	25C	25C22	163N	61.59
25C	25C22	26CD1	53.€5	1610	25C22	162CA	33.51
1610	25C22	163N	60.45	24CA	25C22	26CD1	44.43

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			TAI	BLE XVI			
162CA	25C22	163N	29.15	163N	25C22	26CD1	93.88
25\$G	25023	25N	73.79	25SG	25023	25CB	39.28
25SG	25023	25CA	55.31	25SG	25023	24C	83.85
25SG	25023	190E1	93.34	25SG	25023	162ND1	57.67
25SG	25023	25C	46.54	25SG	25023	26N	46.32
25SG	25023	162CE1	63.52	25N	25023	25CB	50.74
25N	25023	19NE2	75.64	25N	25023	23CA	93.95
25N	25023	23C	70.21	25N	25023	24N	51.30
25N	25023	25CA	24.54	25N	25023	230	71.33
25N	25023	24C	13.19	25N	25023	19CD	72.43
25N	25023	190E1	71.68	25N	25023	24CA	34.54
25N	25023	220	81.27	25N	25023	23N	91.47
25N	25023	25C	27.51	25N	25023	26N	31.41
25N	25023	22C	85.55	25N	25023	162CE1	95.76
25CB	25023	19NE2	84.85	25CB	25023	25CA	26.21
25CB	25023	24C	63.90	25CB	25023	19CD	71.92
25CB	25023	190E1	57.57	25CB	25023	24CA	85.27
25CB	25023	162ND1	56.16	25CB	25023	25C	29.27
25CB	25023	26N	43.07	25CB	25023	162CE1	50.07
19NE2	25023	23CA	79.80	19NE2	25023	23C	86.33
19NE2	25023	24N	70.00	19NE2	25023	25CA	78.56
19NE2	25023	24C	75.95	19NE2	25023	19CD	14.24
19NE2	25023	190E1	31.36	19NE2	25023	24CA	76.27
19NE2	25023	220	39.90	19NE2	25023	162ND1	87.36
19NE2	25023	23N	69.25	19NE2	25023	25C	92.00
19NE2	25023	22C	53.68	19NE2	25023	162CE1	73.60
23CA	25023	23C	26.89	23CA	25023	24N	42.72
23CA	25023	230	37.90	23CA	25023	24C	81.01
23CA .	25023	19CD	93.90	23CA	25023	24CA	59.79
23CA	25023	220	39.98	23CA	25023	23N	10.55
23CA	25023	22C	26.16	23C	25023	24N	23.21
23C	25023	25CA	94.66	23C	25023	230	17.05
23C	25023	24C	57.02	23C	25023	19CD	98.07
23C	25023	24CA	35.84	23C	25023	220	50.35
23C	25023	23N	31.08	23C	25023	25C	92.95
23C	25023	26N	83.85	23C	25023	22C	39.73
24N	25023	25CA	75.53	24N	25023	230	35.32
24N	25023	24C	38.60	24N	25023	19CD	79.21

			TA	BLE XVI			
24N	25023	190E1	91.96	24N	25023	24CA	18.27
24N	25023	220	43.58	24N	25023	23N	40.76
24N	25023	25C	77.49	24N	25023	26N	73.33
24N	25023	22C	39.68	25CA	25023	230	94.04
25CA	25023	24C	37.72	25CA	25023	19CD	69.69
25CA	25023	190E1	61.67	25CA	25023	24CA	59.08
25CA	25023	220	98.54	25CA	25023	162ND1	81.63
25CA	25023	25C	13.70	25CA	25023	26N	29.03
25CA	25023	162CE1	73.11	230	25023	24C	58.97
230	25023	24CA	40.61	230	25023	220	67.32
230	25023	23N	45.18	230	25023	25C	88.46
230	25023	26N	76.05	230	25023	22C	56:12
24C	25023	19CD	76.06	24C	25023	190E1	79.00
24C	25023	24CA	21.38	24C	25023	220	72.85
24C	25023	23N	79.21	24C	25023	25C	39.06
24C	25023	26N	38.39	24C	25023	22C	74.96
19CD	25023	190E1	17.16	19CD	25023	24CA	81.63
19CD	25023	220	53.92	19CD	25023	162ND1	75.34
19CD	25023	23N	83.37	19CD	25023	25C	83.39
19CD	25023	26N	97.86	19CD	25023	22C	67.74
19CD	25023	162CE1	60.88	190E1	25023	24CA	90.13
190E1	25023	220	71.06	190E1	25023	162ND1	60.32
190E1	25023	25C	74.84	190 <b>E1</b>	25023	26N	90.69
190E1	25023	22C	84.88	190 <b>E1</b>	25023	162CE1	45.33
24CA	25023	220	59.13	24CA	25023	23 <b>N</b>	58.89
24CA	25023	25C	59.55	24CA	25023	2 <b>6N</b>	55.16
24CA	25023	22C	57.43	220	25023	23N	29.49
220	25023	22C	13.82	162ND1	25023	25C	84.77
162ND1	25023	26N	95.34	162ND1	25023	162CE1	15.15
23N	25023	22C	15.68	25C	25023	26N	16.15
25C	25023	162CE1	79.30	26N	25023	162CE1	92.61
590	25C24	610D2	91.98	590	25C24	60CA	42.04
590	25C24	61N	69.68	590	25C24		53.63
590	25C24	60C	52.10	590	25C24		9.13
590	25C24	67CD2	94.90	590	25C24		94.67
590	25C24	GON	25.18	590	25C24		98.75
590	25C24	66CA	92.74	590	25C24		94.59
610D2	25C24	60CA	80.40	61002	25C24	61N	53.38

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			TAE	BLE XVI			
610D2	25C24	60C	60.45	610D2	25C24	59C	89.44
610D2	25C24	61CB	30.44	610D2	25C24	60N	85.15
610D2	25C24	650	70.09	610D2	25C24	61CG	12.52
60CA	25C24	61N	33.63	60CA	25C24	60ND2	37.06
60CA	25C24	60C	20.13	60CA	25C24	59C	32.92
60CA	25C24	67CD2	94.92	60CA	25C24	61CB	64.63
60CA	25C24	60N	16.87	60CA	25C24	650	56.77
60CA	25C24	66CA	57.08	60CA	25C24	61CG	74.69
61N	25C24	60ND2	64.13	61N	25C24	60C	17.58
61N	25C24	59C	61.42	61N	25C24	61CB	31.27
61N	25C24	60N	47.40	61N	25C24	650	35.46
61N	25C24	66CA	57.56	61N	25C24	61CG	44.25
60ND2	25C24	60C	56.24	60ND2	25C24	67CE2	75.32
60ND2	25C24	59C	47.84	60ND2	25C24	67CD2	57.88
60ND2	25C24	61CB	93.87	60ND2	25C24	60N	39.84
60ND2	25C24	650	66.16	60ND2	25C24	66CA	42.71
60C	25C24	59C	43.92	60C	25C24	61CB	46.05
60C	25C24	60N	30.63	60C	25C24	650	50.56
60C	25C24	66CA	64.32	60C	25C24	61CG	54.63
67CE2	25C24	67CD2	17.47	67CE2	25C24	66CA	72.97
59C	25C24	67CD2	95.05	5 <b>9</b> C	25C24	61CB	88.03
59C	25C24	60N	16.05	59C	25C24	650	89.64
59C	25C24	66CA	84.79	59C	25C24	61CG	90.22
67CD2	25C24	60N	94.87	67CD2	25C24	650	94.21
67CD2	25C24	66CA	61.75	61CB	25C24	60 <b>N</b>	76.56
61CB	25C24	650	39.70	61CB	25C24	66CA	72.80
61CB	25C24	61CG	18.05	60N	25C24	650	73.63
60N	25C24	66CA	70.86	60 <b>N</b>	25C24	61CG	82.69
650	25C24	66CA	33.53	650	25C24	61CG	57.60
66CA	25C24	61CG	90.84	610D2	25C25	61CB	42.75
610D2	25C25	61N	69.21	61 <b>0D2</b>	25C25	61CG	18.75
610D2	25C25	650	92.92	610 <b>D2</b>	25C25	60C	73.08
610D2	25 <b>C2</b> 5	61CA	51.10	610 <b>D2</b>	25C25	590	97.68
610D2	25C25	60CA	92.69	610D2	25C25	610D1	17.96
610D2	25C25	600	63.26	61CB	25C25	61N	40.79
61CB	25C25	61CG	24.59	61 <b>CB</b>	25C25	650	50.21
61CB	25C25	60C	55.95	61CB	25C25	61CA	21.34
61CB	25C25	60CA	75.07	61CB	25C25	610D1	27.44

	TABLE XVI										
61CB	25C25	65C	60.47	61CB	25C25	600	54.86				
61CB	25C25	66CA	83.99	61N	25C25	61CG	58.00				
61N	25C25	650	42.60	61N	25C25	60C	18.61				
61N	25C25	61CA	20.61	61N	25C25	590	69.42				
61N	25C25	60CA	34.61	61N	25C25	610D1	62.35				
61N	25C25	65C	54.60	61N	25C25	600	25.90				
61N	25C25	66CA	60.64	61CG	25C25	650	74.72				
61CG	25C25	60C	67.35	61CG	25C25	61CA	37.75				
61CG	25C25	60CA	88.04	61CG	25C25	610D1	4.51				
61CG	25C25	65C	84.40	61CG	25C25	600	60.77				
650	25C25	60C	58.12	650	25C25	61CA	47.61				
650	25C25	60CA	61.97	650	25C25	610D1	77.08				
650	25C25	65C	12.55	650	25C25	600	68.17				
650	25C25	66CA	34.76	60C	25C25	61CA	34.62				
60C	25C25	590	51.04	60C	25C25	60CA	20.69				
60C	25C25	610D1	71.85	60C	25C25	65C	68.94				
60C	25C25	600	12.51	60C	25C25	66CA	66.29				
61CA	25C25	590	84.06	61CA	25C25	60CA	53.97				
61CA	25C25	610D1	41.99	61CA	25C25	65C	60.01				
61CA	25C25	600	34.68	61CA	25C25	66CA	75.36				
590	25C25	60CA	38.70	590	25C25	600	49.63				
590	25C25	66CA	83.86	60CA	25C25	610D1	92.54				
60CA	25C25	65C	69.48	60CA	25C25	600	29.48				
60CA	25C25	66CA	56.22	610D1	25C25	65C	86.17				
610D1	25C25	600	65.14	65C	25C25	600	79.72				
65C	25C25	66CA	29.00	600	25C25	66CA	78.80				
61CB	25C26	650	57.98	61CB	25C26	610D2	40.43				
61CB	25C26	61CG	23.23	61CB	25C26	61N	39.07				
61CB	25C26	65C	73.35	61CB	25C26	61CA	18.97				
61CB	25C26	640	64.81	61CB	25C26	66N	88.64				
61CB	25C26	66CA	94.09	61CB	25C26	60C	49.39				
61CB	25C26	65 <b>CA</b>	72.50	61CB	25C26	64C	54.69				
61CB	25C26	610D1	25.08	61CB	25C26	60CA	66.90				
650	25C26	610D2	96.74	650	25C26	61CG	81.20				
650	25C26	61N	45.23	630	25C26	65C	17.69				
650	25C26	61CA	49.81	650	25C26	640	56.41				
650	25C26		30.68	650	25C26	66CA	39.57				
650	25C26	60C	55.87	650	25C26	65CA	28.89				

	TABLE XVI									
650	25C26	64C	44.41	650	25C26	610D1	82.28			
650	25C26	60CA	58.71	610D2	25C26	61CG	19.08			
610D2	25C26	61N	60.79	610D2	25C26	61CA	47.71			
610D2	25C26	640	94.75	610D2	25C26	60C	60.86			
610D2	25C26	64C	89.54	610D2	25C26	610D1	23.61			
610D2	25C26	60CA	75.67	61CG	25C26	61N	54.13			
61CG	25C26	65C	96.40	61CG	25C26	61CA	36.24			
61CG	25C26	640	76.97	61CG	25C26	60C	59.38			
61CG	25C26	65CA	93.65	61CG	25C26	64C	70.68			
61CG	25C26	610D1	7.62	61CG	25C26	60CA	76.74			
61N	25C26	65C	62.37	61N	25C26	61CA	20.12			
61N	25C26	640	88.15	61N	25C26	66N	69.41			
61N	25C26	66CA	64.66	61N	25C26	60C	13.98			
61N	25C26	65CA	73.06	61N	25C26	64C	74.50			
61N	25C26	610D1	59.89	61N	25C26	60CA	28.83			
65C	25C26	61CA	67.25	65C	25C26	640	51.79			
65C	25C26	6 <b>6N</b>	17.38	65C	25C26	66CA	33.55			
65C	25C26	60C	71.82	65C	25C26	65CA	17.23			
65C	25C26	64C	43.45	65C	25C26	610D1	96.15			
65C	25C26	60CA	71.11	61CA	25C26	640	76.04			
61CA	25C26	6 <b>6N</b>	79.28	61CA	25C26	66CA	79.89			
61CA	25C26	60C	31.02	61CA	25C26	65CA	72.24			
61CA	25C26	64C	63.54	61CA	25C26	610D1	40.94			
61CA	25C26	60CA	48.15	640	25C26	66N	65.51			
640	25C26	66CA	84.31	640	25C26	65CA	35.23			
640	25C26	64C	13.78	640	25C26	610D1	71.21			
6 <b>6</b> N	25C26	66CA	19.09	6 <b>6N</b>	25C26	60C	75.28			
6 <b>6N</b>	25C26	65CA	30.56	6 <b>6N</b>	25C26	64C	59.39			
66N	25C26	60CA	69.05	56CA	25C26	60C	66.08			
66CA	25C26	65CA	49.11	6 <b>6CA</b>	25C26	64C	76.99			
66CA	25C26	60CA	55.34	60C	25C26	65CA	84.60			
60C	25C26	64C	88.43	60C	25C26	610D1	66.22			
60C	25C26	60CA	17.82	65CA	25C26	64C	29.29			
65CA	25C26	610D1	91.21	65CA	25C26	60CA	86.87			
64C	25C26	610D1	66.22	64C	25C25	60CA	98.77			
610D1	25C26	60 <b>C</b> A	83.78	650	25C 7	65C	15.69			
650	25C27	66N	35.95	650	25C17	66CA	44.75			
650	25C27	65CA	32.29	650	25C27	640	53.20			

	TABLE XVI								
650	25C27	61CB	45.32	650	25C27	66C	62.38		
650	25C27	61N	36.43	650	25C27	610D2	72.86		
650	25C27	660	66.33	65C	25C27	66N	20.59		
65C	25C27	66CA	38.16	65C	25C27	65CA	18.27		
65C	25C27	640	50.92	65C	25C27	61CB	62.98		
65C	25C27	66C	52.26	65C	25C27	61N	55.57		
65C	25C27	610D2	91.46	65C	25C27	660	51.89		
66N	25C27	66CA	22.16	66N	25C27	65CA	32.66		
66N	25C27	640	68.31	66N	25C27	61CB	81.26		
66N	25C27	66C	32.35	66N	25C27	61N	66.24		
66N	25C27	67CD2	86.04	66N	25C27	660	31.43		
66CA	25C27	67CE2	84.05	66CA	25C27	65CA	53.89		
66CA	25C27	640	88.87	66CA	25C27	61CB	86.12		
66CA	25C27	66C	18.33	66CA	25C27	61N	61.78		
66CA	25C27	67CD2	68.00	66CA	25C27	67CZ	89.01		
66CA	25C27	660	27.54	67CE2	25C27	66C	68.52		
67CE2	25C27	67CD2	16.65	67CE2	25C27	67CZ	16.31		
67CE2	25C27	660	70.63	65CA	25C27	640	35.68		
65CA	25C27	61CB	65.69	65CA	25C27	66C	64.52		
65CA	25C27	61N	68.03	65CA	25C27	610D2	94.09		
65CA	25C27	660	59.86	640	25C27	61CB	54.95		
640	25C27	61N	76.01	640	25C27	610D2	75.54		
640	25C27	660	93.93	61CB	25C27	61N	30.76		
61CB	25C27	610D2	28.82	66C	25C27	61N	78.83		
66C	25C27	67CD2	53.68	66C	25C27	67CZ	71.38		
66C	25C27	6 <b>6</b> 0	13.90	61N	25C27	610D2	45.79		
61N	25C27	660	89.25	67CD2	25C27	67CZ	28.76		
67CD2	25C27	660	58.30	67CZ	25C27	660	69.31		
67CE2	`25C28	67CD2	23.52	67CE2	25C28	67CZ	20.96		
67CE2	25C28	670H	35.20	67C <b>E2</b>	25C28	66C	86.92		
67CE2	25C28	67CG	33.10	67CE2	25C28	67N	73.57		
67CE2	25C28	67CE1	29.88	67CE2	25C28		85.55		
67CE2	25C28	660	85.99	67CE2			33.21		
67CD2	25C28	67CZ	38.76	67CD2	25C28	66CA	84.93		
67CD2	25C28	670H	56.87	67CD2	25C28	66C	66.69		
67CD2	25C28	67CG	16.19	67CD2	25C28		51.42		
67CD2	25C28	67CE1	37.73	67CD2			63.64		
67CD2	25C28	660	69.45	67CD2	25C28	67CD1	28.35		

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			TAP	BLE XVI			•
67CZ	25C28	670H	19.51	67CZ	25C28	66C	88.11
67CZ	25C28	67CG	39.85	67CZ	25C28	67N	79.74
67CZ	25C28	67CE1	15.82	67CZ	25C28	660	82.16
67CZ	25C28	67CD1	29.69	66CA	25C28	66C	21.36
66CA	25C28	66N	20.18	66CA	25C28	67CG	73.95
66CA	25C28	650	39.39	66CA	25C28	67N	33.53
66CA	25C28	65C	33.20	66CA	25C28	67CE1	94.77
66CA	25C28	60ND2	46.84	66CA	25C28	660	30.13
66CA	25C28	67CD1	79.87	670H	25C28	67CG	59.35
670H	25C28	67N	98.79	670H	25C28	67CE1	31.87
670H	25C28	660	96.91	670H	25C28	67CD1	48.13
66C	25C28	66N	33.87	66C	25C28	67CG	53.83
66C	25C28	650	60.02	66C	25C28	67N	17.65
66C	25C28	65C	50.69	66C	25C28	67CE1	73.54
66C	25C28	60ND2	49.87	66C	25C28	660	14.26
66C	25C28	67CD1	58.53	66N	25C28	67CG	86.97
66N	25C28	650	30.60	66N	25C28	67N	50.32
66N	25C28	65C	17.37	6 <b>6N</b>	25C28	67CE1	99.34
66N	25C28	60ND2	66.19	66N	25C28	660	33.91
66N	25C28	67CD1	87.99	67CG	25C28	67N	41.20
67CG	25C28	67CE1	31.84	67CG	25C28	60ND2	64.79
67CG	25C28	660	54.35	67CG	25C28	67CD1	16.69
650	25C28	67N	72.54	650	25C28	65C	16.08
650	25C28	60ND2	67.97	650	25C28	660	63.89
67N	25C28	65C	66.13	67N	25C28	67CE1	67.52
67N	25C28	60ND2	39.36	67N	25C28	660	28.39
67N	25C28	67CD1	50.73	65C	25C28	60ND2	73.05
65C	25C28	660	51.14	67CE1	25C28	60ND2	96.62
67CE1	25C28	660	66.57	67CE1	25C28	67CD1	17.05
60ND2	25C28	660		60ND2	25C28		
660	25C28	67CD1	54.11	67CE2	25C29	67CD2	24.48
67CE2	25C29	60ND2	95.45				15.31
67CE2	25C29	66CA	94.75	67CE2	25C29	67CG	30.54
67CE2	25 <b>C29</b>	670H	29.52	67CE2	25C29	67N	69.16
67CE2	25C29	66C	76.62	67CE2	25C29	700D1	69.19
67CD2	25C29	60ND2	70.39	67CD2	25C29	67CZ	35.94
67CD2	25C29	66CA	77.68	67CD2	25C29		11.52
67CD2	25C29	670H	52.86	67CD2	25C29	67N	48.30

TABLE XVI							
67CD2	25C29	66C	59.66	67CD2	25C29	700D1	48.35
60ND2	25C29	590	51.92	60ND2	25C29	66CA	47.48
60ND2	25C29	67CG	67.53	60ND2	25C29	67N	39.60
60ND2	25C29	60CA	34.92	60ND2	25C29	66C	48.58
60ND2	25C29	700D1	36.76	60ND2	25C29	650	66.66
590	25C29	66CA	91.68	590	25C29	67N	91.44
590	25C29	60CA	36.13	590	25C29	66C	99.32
590	25C29	700D1	57.47	590	25C29	650	87.31
67CZ	25C29	66CA	92.71	67CZ	25C29	67CG	37.70
67CZ	25C29	670H	17.56	67CZ	25C29	67N	72.63
67CZ	25C29	66C	75.91	67CZ	25C29	700D1	83.35
66CA	25C29	67CG	67.01	66CA	25C29	67N	30.81
66CA	25C29	60CA	57.52	66CA	25C29	66C	18.31
66CA	25C29	700D1	76.75	66CA	25C29	650	34.30
67CG	25C29	670H	55.25	67CG	25C29	67 <b>N</b>	38.92
67CG	25C29	66C	48.79	67CG	25C29	700D1	52.17
670H	25C29	67N	88.91	670H	25C29	66C	89.71
670H	25C29	700D1	98.59	67N	25C29	60CA	68.41
67N	25C29	66C	16.09	67N	25C29	700D1	52.40
67N	25C29	650	64.97	60CA	25C29	66C	69.66
60CA	25C29	700D1	64.85	60CA	25C29	650	52.85
66C	25C29	700D1	68.06	66C	25C29	650	51.98
65C	25C30	66N	22.36	65C	25C30	650	19.60
65C	25C30	65CA	24.11	65C	25C30	640	60.69
65C	25C30	66CA	37.47	65C	25C30	64C	47.70
65C	25C30	65N	31.97	65C	25C30	66C	52.44
65C	25C30	660	56.13	6 <b>6N</b>	25C30	650	36.76
66N	25C30	65CA	39.03	66 <b>N</b>	25C30	640	81.12
6 <b>6N</b>	25C30	66CA	20.96	6 <b>6N</b>	25C30	64C	68.55
6 <b>6</b> N	25C30	65N	51.94	66N	25C30	66C	31.15
66N	25C30	660	33.85	650	25C30	65CA	37.88
650	25C30	640	60.09	650	25C30	66CA	42.53
650	25C30	64C	47.65	650	25C30	65N	37.00
650	25C30	66C	60.78	650	25C30	660	69.10
65CA	25C30	640	43.45	65CA	25C30	66CA	58.77
65CA	25C30	64C	32.33	65CA	25C30	65N	16.26
65CA	25C30	66C	69.64	65CA	25C30	660	67.50
640	25C30	66CA	98.08	540	25C30	64C	13.08

			TAE	LE XVI			
640	25C30	65N	29.19	66CA	25C30	64C	85.02
66CA	25C30	65N	69.41	66CA	25C30	66C	18.61
66CA	25C30	660	30.64	64C	25C30	65N	16.80
64C	25C30	66C	99.65	64C	25C30	660	99.64
65N	25C30	66C	83.08	65N	25C30	660	82.93
66C	25C30	660	15.25	66N	25031	65C	17.32
66N	25031	67CZ	97.91	66N	25031	65CA	31.08
66N	25031	640	62.95	6 <b>6N</b>	25031	66CA	17.26
66N	25031	660	33.20	66N	25031	67CE1	92.81
66N	25031	67CE2	86.83	66N	25031	650	26.88
66N	25031	66C	29.04	65C	25031	65CA	19.09
65C	25 <u>0</u> 31	640	46.34	65C	25031	66CA	30.23
65C	25031	660	50.48	65C	25031	67CE2	97.89
65C	25031	650	13.61	65C	25031	66C	45.49
670H	25031	67CZ	17.55	670H	25031	66CA	98.18
670H	25031	660	90.84	670H	25031	67CE1	29.52
670H	25031	67CE2	29.04	670H	25031	66C	87.47
67CZ	25031	66CA	81.60	67CZ	25031	660	73.49
67CZ	25031	67CE1	16.52	67CZ	25031	67CE2	16.35
67CZ	25031	66C	69.99	65CA	25031	640	34.51
65CA	25031	66CA	47.38	65CA	25031	660	60.33
65CA	25031	650	28.52	65CA	25031	66C	59.78
640	25031	66CA	76.15	640	25031	660	94.57
640	25031	650	44.97	640	25031	66C	91.77
66CA	25031	660	28.82	66CA	25031	67CE1	78.69
66CA	25031	67CE2	69.71	66CA	25031	650	33.12
66CA	25031	66C	17.65	660	25031	67CE1	63.44
660	25031	67CE2	68.73	660	25031	650	58.85
660	25031	66C	14.31	67CE1	25031	67CE2	28.45
67CE1	25031	66C	63.77	57CE2	25031	650	93.08
67CE2	25031	66C	61.53	650	25031	66C	50.53
660	25 <b>C</b> 32	66N	36.46	660	25C32	65CA	64.42
660	25C32	65C	52.71	660	25C32	67CE1	69.47
660	25C32	66C	14.32	660	25C32	67CZ	76.69
660	25C32	66CA	29.45	660	25C32	670H	92.97
66N	25C32	65CA	31.01	6611	25C32	65C	16.26
66N	25C32	67CE1	96.14	6 <b>6N</b>	25C32	66C	30.68
66N	25C32	67CZ	95.21	6 G N	25C32	66CA	16.64

TABLE XVI									
65CA	25C32	65C	18.68	65CA	25C32	66C	61.49		
65CA	25C32	66CA	46.86	65C	25C32	66C	46.16		
65C	25C32	66CA	29.55	67CE1	25C32	6 <b>6</b> C	67.09		
67CE1	25C32	67CZ	16.76	67CE1	25C32	66CA	79.83		
67CE1	25C32	67OH	29.00	66C	25C32	67CZ	70.31		
66C	25C32	66CA	18.10	66C	25C32	670H	85.85		
67CZ	25C32	66CA	78.64	67CZ	25C32	670H	16.35		
66CA	25C32	670H	92.05	67CE1	25033	67CZ	18.11		
67CE1	25033	670H	32.18	67CE1	25033	660	70.79		
67CE1	25033	67CD1	16.55	67CE1	25033	2530H2	91.77		
67CZ	25033	670H	17.82	67CZ	25033	660	76.31		
67CZ	25033	67CD1	30.29	67CZ	25033	2530H2	94.53		
670H	25033	660	93.31	670H	25033	67CD1	47.11		
670H	25033	253OH2	82.17	660	25033	67CD1	55.11		
2530H2	25033	1600	32.53	660	25C34	6 <b>6N</b>	39.24		
660	25C34	65CA	72.10	660	25C34	65C	54.84		
660	25C34	25SG	98.30	660	25C34	66C	10.30		
66N	25C34	65CA	33.35	6 <b>6N</b>	25C34	65C	15.62		
66N	25C34	25SG	96.34	66N	25C34	66C	31.01		
65CA	25C34	65C	18.75	65CA	25C34	25 <b>S</b> G	85.74		
65CA	25C34	66C	64.35	65C	25C34	25 <b>S</b> G	95.57		
65C	25C34	66C	46.29	25SG	25C34	1610	60.31		
25 <i>S</i> G	25C34	161C	66.90	1610	25C34	161C	14.93		
660	25C35	66C	7.49	660	25C35	68CE	57.46		
660	25C35	66N	35.66	660	25C35	163CB	89.86		
660	25C35	66CA	21.29	6 <b>6</b> C	25C35	68CE	64.25		
66C	25C35	66N	32.33	66C	25C35	163CB	97.34		
66C	25C35	66CA	16.42	68CE	25C35	66N	88.65		
68CE	25C35	163CB	42.93	SSCE	25C35		71.03		
68CE	25C35	66CA	78.09	6 <b>6N</b>	25C35				
163CB	25C35	163N	28.84	660	25C36				
660	25C36	67CD1	56.25	660	25 <b>C</b> 36	67CE1			
134CB	23C36	209CD2	53.33	134CB			70.27		
134CB	25C36	1600	87.04	134CB		162N	72.66		
134CB	25C36	160C	74.39	134CB		161C	86.84		
134CB	25C36	161CA	88.42	134CB		161N	74.11		
134CB	25C36	160CB	54.87	209CD2		68CE	80.58		
209CD2	25C33	1600	88.38	209CD2	25C36	160C	87.02		

			TA	BLE XVI			
209CD2	25C36	67CD1	54.52	209CD2	25C36	67CE1	55.35
209CD2	25C36	161N	98.98	209CD2	25C36	160CB	57.71
68CE	25C36	67CD1	70.75	68CE	25C36	67CE1	87.36
1600	25C36	162N	60.06	1600	25C36	160C	14.84
1600	25C36	161C	52.10	1600	25C36	161CA	34.02
1600	25C36	161N	27.16	1600	25C36	160CB	35.85
162N	25C36	160C	49.38	162N	25C36	161C	15.90
162N	25C36	161CA	29.60	162N	25C36	161N	33.75
162N	25C36	160CB	69.36	160C	25C36	161C	45.30
160C	25C36	161CA	29.33	160C	25C36	161N	15.76
160C	25C36	160CB	29.68	161C	25C36	161CA	18.21
161C	25C36	161N	30.25	161C	25C36	160CB	71.28
161CA	25C36	161N	17.23	161CA	25C36	160CB	58.05
67CD1	25C36	67CE1	16.61	67CE1	25C36	160CB	96.33
161N	25C36	160CB	41.59	162N	25C37	162C	38.77
162N	25C37	163N	55.65	162N	25C37	1620	43.88
162N	25C37	161C	19.66	162N	25C37	161N	43.11
162N	25C37	162CA	20.61	162N	25C37	161CA	36.42
162N	25C37	160C	61.64	162N	25C37	163CA	72.52
162N	25C37	163CB	88.83	162N	25C37	1600	71.57
162N	25C37	160CB	86.86	162N	25C37	134CA	86.74
162N	25C37	1610	27.40	162N	25C37	160CA	69.38
134CB	25C37	162C	80.90	134CB	25C37	163N	83.96
134CB	25C37	1620	64.46	134CB	25C37	161N	96.15
134CB	25C37	162CA	99.27	134CB	25C37	160C	92.07
134CB	25C37	163CA	69.42	134CB	25C37	163CB	75.64
134CB	25C37	160CB	64.54	134CB	25C37	134CA	17.37
134CB	25C37	160CA	74.76	134CB	25C37	68CE	75.20
134CB	25C37	209CD2	53.65	162C	25C37	163N	20.04
162C	25C37	1620	18.39	162C	25C37	161C	56.63
162C	25C37	161N	77.75	162C	25C37	162CA	22.08
1.62C	25C37	161CA	75.17	16 <b>2C</b>		160C	94.90
152C	25C37	163CA	33.93	162C		163CB	51.74
162C	25C37	134CA	63.54	162C	25C37	1610	57.80
162C	25C37	160CA	95.14	162C	25C37	68CE	93.53
163N	25C37	1620	33.61	163N	25C37	161C	70.64
163N	25C37	161N	97.07	163N		162CA	
163N	25C37	161CA	91.41	1.63N	25C37	163CA	19.33

			TA	BLE XVI			
163N	25C37	163CB	33.19	163N	25C37	134CA	67.66
163N	25C37	1610	67.69	163N	25C37	68CE	75.10
1620	25C37	161C	63.53	1620	25C37	161N	73.01
1620	25C37	162CA	34.99	1620	25C37	161CA	77.20
1620	25C37	160C	87.26	1620	25C37	163CA	38.61
1620	25C37	163CB	58.25	1620	25C37	160CB	91.83
1620	25C37	134CA	47.36	1620	25C37	1610	68.76
1620	25C37	160CA	82.84	1620	25C37	68CE	96.46
161C	25C37	161N	37.46	161C	25C37	162CA	35.43
161C	25C37	161CA	21.87	161C	25C37	160C	54.49
161C	25C37	163CA	89.14	161C	25C37	1600	59.67
161C	25C37	160CB	85.75	161C	25C37	1610	13.22
161C	25C37	160CA	67.48	161N	25C37	162CA	63.61
161N	25C37	161CA	20.70	161N	25C37	160C	18.54
161N	25C37	1600	30.38	161N	25C37	160CB	48.33
161N	25C37	134CA	92.03	161N	25C37	1610	49.54
161N	25C37	160CA	30.13	162CA	25C37	161CA	55.77
162CA	25C37	160C	82.09	162CA	25C37	163CA	53.72
162CA	25C37	163CB	68.64	162CA	25C37	1600	92.06
162CA	25C37	134CA	82.32	162CA	25C37	1610	35.74
162CA	25C37	160CA	88.24	161CA	25C37	160C	34.21
161CA	25C37	1600	37.80	161CA	25C37	160CB	67.22
161CA	25C37	1610	31.20	161CA	25C37	160CA	49.65
160C	25C37	1600	15.98	160C	25C37	160CB	33.38
160C	25C37	134CA	93.50	160C	25C37	1610	65.32
160C	25C37	160CA	17.61	160C	25C37	209CD2	89.23
163CA	25C37	163CB	19.64	163CA	25C37	134CA	55.20
163CA	25C37	1610	86.98	163CA	25C37	68CE	59.96
163CB	25C37	134CA	65.55	163CB	25C37	1610	97.16
163CB	25C37	68CE	41.99	1600	25C37	160CB	38.98
1600	25C37	1610	67.69	1600	25C37	160CA	29.34
1600	25C37	209CD2	86.31	160CB	25C37	134CA	72.65
160CB	25C37	1610	97.76	160CB	25C37	160CA	18.30
160CB	25C37	209CD2	58.20	134CA	25C37	160CA	77.94
134CA	25C37	68CE	77.56	134CA		209CD2	71.01
1610	25C37	160CA	79.66	160CA	25C37	209CD2	76.48
68CE	25C37	209CD2	74.72	209CD2	25C38	67CD1	74.87
209CD2	25C38	67CE1	73.16	209CD2	25C38	134CB	65.56

			TA	BLE XVI			
209CD2	25C38	209CG	16.73	209CD2	25C38	67CG	84.81
67CD1	25C38	67CE1	20.70	67CD1	25C38	68CE	89.96
67CD1	25C38	660	66.53	67CD1	25C38	209CG	75.95
67CD1	25C38	67CG	12.21	67CD1	25C38	67CA	42.12
67CE1	25C38	660	77.82	67CE1	25C38	209CG	80.22
67CE1	25C38	67CG	30.41	67CE1	25C38	67CA	60.82
68CE	25C38	134CB	80.62	68CE	25C38	660	53.63
68CE	25C38	209CG	90.75	68CE	25C38	67CG	80.08
68CE	25C38	67CA	50.64	134CB	25C38	209CG	59.36
660	25C38	67CG	54.73	660	25C38	67CA	34.43
209CG	25C38	67CG	83.39	209CG	25C38	67 <b>CA</b>	92.38
67CG	25C38	67CA	30.58	65CA	25C39	66N	39.58
65CA	25C39	660	81.69	65CA	25C39	26CD1	58.77
65CA	25C39	65C	21.13	65CA	25C39	26CB	87.77
65CA	25C39	26CG	70.14	65CA	25C39	66C	70.43
65CA	25C39	65N	10.50	65CA	25C39	26N	97.84
65CA	25C39	66CA	51.95	65CA	25C39	230	44.24
25SG	25C39	26CD1	80.82	25 <b>S</b> G	25C39	26CB	77.99
25SG	25C39	26CG	82.69	25 <i>S</i> G	25C39	26N	48.23
25SG	25C39	230	74.80	25 <i>S</i> G	25C39	1610	65.02
6 6 N	25C39	660	42.13	6 <b>6N</b>	25C39	26CD1	49.89
66N	25C39	65C	18.53	6 <b>6N</b>	25C39	26CB	61.31
66N	25C39	26CG	50.14	6 <b>6N</b>	25C39	66C	30.88
66N	25C39	65N	49.74	6 <b>6N</b>	25C39	26N	85.92
66N	25C39	66CA	12.40	6 <b>6N</b>	25C39	230	70.96
660	25C39	26CD1	64.20	660	25C39	65C	60.58
660	25C39	26CB	46.66	660	25C39	26CG	51.76
660	25C39	66C	11.26	660	25C39	65N	91.86
660	25C39	26N	76.72	660	25C39	66CA	29.74
26CD1	25C39	65C	52.77	26CD1	25C39	26CB	33.76
26CD1	25C39	26CG	16.02	26CD1	25C39	66C	59.08
26CD1	25C39	65 <b>N</b>	61.27	26CD1	25C39	26N	40.34
26CD1	25C39	66CA	52.43	26CD1	25C39	230	46.03
65C	25C39	26CB	74.04	65C	25C39	26CG	59.01
65C	25C39	66C	49.32	65C	25C39	65N	31.45
65C	25C39	26N	92.79	65C	25C39	66CA	30.85
65C	25C39	230	58.16	26CB	25C39	26CG	18.51
2.6CB	25C39	66C	48.81	26CB	25C39	65N	92.88

TABLE XVI							
26CB	25C39	26N	30.42	26CB	25C39	66CA	55.33
26CB	25C39	230	78.36	26CG	25C39	66C	49.18
26CG	25C39	65N	74.55	26CG	25C39	26N	35.79
26CG	25C39	66CA	48.23	26CG	25C39	230	61.96
66C	25C39	65N	80.62	66C	25C39	26N	79.15
66C	25C39	66CA	18.48	66C	25C39	230	96.37
65N	25C39	26N	97.76	65N	25C39	66CA	62.14
65N	25C39	230	37.19	26N	25C39	66CA	83.19
26N	25C3 <sub>.</sub> 9	230	67.94	66CA	25C39	230	81.21
66N	25040	660	55.03	66N	25040	26CD1	68.97
66N	25040	65CA	46.68	66N	25040	26CB	85.08
66N	25040	26CG	70.07	66N	25040	65C	21.06
66N	25040	66C	40.07	66N	25040	66CA	17.56
6 <b>6</b> N	25040	26NE1	58.94	66N	25040	65N	53.77
66N	25040	230	82.65	66N	25040	26CD2	60.09
66N	25040	650	19.20	66N	25040	26CE2	54.44
660	25040	26CD1	89.64	660	25040	26CB	63.20
660	25040	26CG	71,47	660	25040	65C	75.92
660	25040	66C	15.74	660	25040	66CA	38.45
660	25040	26N	99.32	660	25040	26NE1	91.06
660	25040	26CA	79.34	660	25040	26CD2	68.55
660	25040	650	74.21	660	25040	68CE	47.85
660	25040	26CE2	78.85	26CD1	25040	65CA	74.34
26CD1	25040	26CB	46.12	26CD1	25040	26CG	22.72
26CD1	25040	65C	67.94	26CD1	25040	66C	79.23
26CD1	25040	66CA	69.49	26CD1	25040	25 <b>S</b> G	92.07
26CD1	25040	26N	50.30	26CD1	25040	26NE1	14.37
26CD1	25040	26CA	50.84	26CD1	25040	65N	70.61
26CD1	25040	230	51.41	26CD1	25040	26CD2	21.35
26CD1	25040	650	66.27	26CD1	25040	68CE	92.41
26CD1	25040	26CE2	14.76	26CD1	25040	25N	52.65
65CA	25040	26CG	91.43	65CA	25040	65C	25.75
55CA	25040	66C	86.58	65CA	25040	66CA	64.21
65CA	25040	26NE1	60.05	65CA	25040	65N	8.97
65CA	25040	230	48.18	65CA	25040	26CD2	82.47
65CA	25040	650	27.92	65CA	25040	26CE2	67.12
65CA	25040	25N	97.61	26CB	25040	26CG	24.84
26CB	25040	65C	97.95	26CB	25040	66C	63.€2

			TAE	BLE XVI			
26CB	25040	66CA	72.60	26CB	25040	25SG	89.79
26CB	25040	26N	37.46	26CB	25040	26NE1	58.62
26CB	25040	26CA	19.07	26CB	25040	230	95.25
26CB	25040	26CD2	33.11	26CB	25040	650	95.49
26CB	25040	68CE	46.43	26CB	25040	26CE2	49.00
26CB	25040	25N	67.75	26CG	25040	65C	77.41
26CG	25040	66C	64.77	26CG	25040	66CA	63.32
26CG	25040	25 <i>S</i> G	97.02	26CG	25040	26N	44.89
26CG	25040	26NE1	33.94	26CG	25040	26CA	35.84
26CG	25040	65N	89.76	26CG	25040	230	74.03
26CG	25040	26CD2	10.27	26CG	25040	650	75.16
26CG	25040	68CE	70.18	26CG	25040	26CE2	24.36
26CG	25040	25N	62.89	65C	25040	66C	61.12
65C	25040	66CA	38.51	65C	25040	26NE1	54.62
65C	25040	65N	32.73	65C	25040	230	65.00
65C	25040	26CD2	67.24	65C	25040	650	2.52
65C	25040	26CE2	55.73	66C	25040	66CA	22.95
66C	25040	26NE1	78.03	66C	25040	26CA	81.98
66C	25040	65 <b>N</b>	93.83	66C	25040	26CD2	59.43
66C	25040	650	59.25	66C	25040	68CE	60.81
66C	25040	26CE2	66.69	66CA	25040	26NE1	63.40
66CA	25040	26CA	91.57	66CA	25040	65N	71.07
66CA	25040	230	96.11	66CA	25040	26CD2	54.65
66CA	25040	650	36.50	66CA	25040	68CE	82.71
66CA	25040	26CE2	54.95	25 <b>S</b> G	25040	26N	53.76
25SG	25040	26CA	70.74	25 <b>S</b> G	25040	230	76.38
25SG	25040	68CE	91.56	25 <b>S</b> G	25040	25N	39.87
26N	25040	26NE1	64.06	2 <b>6N</b>	25040	26CA	19.99
26N	25040	230	77.54	26N	25040	26CD2	54.76
26N	25040	68CE	65.32	26N	25040	26CE2	62.82
26N	25040	25 <b>N</b>	33.74	26 <b>NE1</b>	25040	26CA	65.10
26NE1	25040	65N	56.73	26 <b>NE</b> 1	25040	230	44.12
26NE1	25040	26CD2	28.39	26NE1	25040	650	53.18
26NE1	25040	26CE2	12.89	26NE1	25040	25N	60.41
26CA	25040	230	91.40	26CA	25040	26CD2	45.97
26CA	25040	68CE	49.24	26CA	25040	26CE2	59.06
26CA	25040	25N	53.14	65 <b>%</b>	25043	230	39.46
65N	25040	26CD2	81.69	65 <b>N</b>	2504	650	34.59

	TABLE XVI								
65N	25040	26CE2	65.52	65N	25040	25N	89.01		
230	25040	26CD2	71.57	230	25040	650	65.65		
230	25040	26CE2	57.01	230	25040	25N	49.58		
26CD2	25040	650	64.96	26CD2	25040	68CE	75.92		
26CD2	25040	26CE2	16.58	26CD2	25040	25N	69.33		
650	25040	26CE2	53.78	68CE	25040	26CE2	92.46		
68CE	25040	25N	97.92	26CE2	25040	25N	67.27		
25SG	25N41	1610	80.59	25SG	25N41	230	90.56		
25SG	25N41	26CD1	82.77	25 <i>S</i> G	25N41	25CB	9.91		
25SG	25N41	23C	79.09	25SG	25N41	25N	40.23		
25SG	25N41	161C	80.64	25SG	25N41	26N	46.58		
25SG	25N41	162CA	52.67	25SG	25N41	163N	39.29		
65CA	25N41	230	48.79	65CA	25N41	26CD1	53.27		
65CA	25N41	66N	31.72	65CA	25N41	23C	63.36		
65CA	25N41	65N	14.93	65CA	25 <b>N4</b> 1	65C	15.53		
65CA	25N41	25N	93.22	65CA	25N41	26N	91.88		
1610	25 <b>N4</b> 1	25CB	89.53	1610	25N41	161C	13.63		
1610	25N41	162CA	33.89	1610	25N41	163N	62.20		
230	25N41	26CD1	48.03	230	25 <b>N4</b> 1	6 <b>6N</b>	68.57		
230	25N41	25CB	80.86	230	25N41	23C	14.88		
230	25N41	65N	39.21	230	25N41	65C	57.67		
230	25N41	25N	51.81	230	25N41	26N	71.06		
26CD1	25N41	6 <b>6N</b>	42.24	26CD1	25N41	25CB	74.76		
26CD1	25N41	23C	56.38	26CD1	25N41	6 <b>5N</b>	58.93		
26CD1	25N41	65C	46.09	26CD1	25N41	25N	50.77		
26CD1	25N41	26N	38.61	26CD1	25 <b>N4</b> 1	163N	98.70		
66N	25N41	23C	82.86	6 <b>6N</b>	25N41	65N	46.23		
66N	25N41	65C	16.19	6 <b>6N</b>	25 <b>N4</b> 1	25N	92.91		
66N	25N41	26N	75.02	25CB	25N41	23C	69.90		
25CB	25N41	25N	30.33	25CB	25N41	161C	90.33		
25CB	25 <b>N4</b> 1	26N	40.90	25CB	25N41	162CA	62.56		
25CB	25N41	163N	48.01	23C	25N41	65N	52.61		
23C	25N41	65C	72.52	23C	25N41	25N	43.58		
23C	25N41	26N	69.77	65 <b>N</b>	25N41	65C	30.16		
65N	25N41	25N	89.00	65N	25N41	26N	96.17		
65C	25N41	25N	93.25	65C	25N41		83.49		
25N	25N41	26N	33.34	25N		162CA			
25N	25N41	163N	75.53	161C	25N41	162CA	28.77		

			TA	BLE XVI			•
161C	25N41	163N	54.46	26N	25N41	162CA	89.00
26N	25N41	163N	62.08	162CA	25N41	163N	28.61
25 <i>S</i> G	25C42	25N	61.47	25SG	25C42	25CB	23.02
25SG	25C42	24N	99.07	25 <b>S</b> G	25C42	26N	58.96
25SG	25C42	25CA	42.76	25SG	25C42	24C	76.26
25 <b>S</b> G	25C42	24CA	95.01	25SG	25C42	25C	46.56
25 <b>S</b> G	25C42	26CG	94.70	25SG	25C42	26CB	80.63
230	25C42	23C	20.90	230	25C42	25N	72.70
230	25C42	26CD1	61.41	230	25C42	65CA	57.62
230	25C42	23CA	37.29	230	25C42	24N	34.67
230	25C42	26N	92.67	230	25C42	25CA	92.41
230	25C42	24C	59.75	230	25C42	24CA	39.73
230	25C42	65 <b>N</b>	43.51	230	25C42	26NE1	46.24
230	25C42	25C	96.61	230	25C42	26CG	73.49
230	25C42	6 6 N	75.31	230	25C42	65C	62.99
230	25C42	26CB	90.54	23C	25C42	25N	60.95
23C	25C42	26CD1	74.34	23C	25C42	65CA	77.95
23C	25C42	25CB	94.36	23C	25C42	23CA	22.31
23C	25C42	24N	19.02	23C	25C42	2 <b>6N</b>	92.21
23C	25C42	25CA	80.52	23C	25C42	24C	52.38
23C	25C42	24CA	33.67	23C	25C42	65N	62.45
23C	25C42	26NE1	61.04	23C	25C42	25C	90.42
23C	25C42	26CG	85.04	23C	25C42	66N	95.80
23C	25C42	65C	83.89	25N	25C42	26CD1	66.35
25N	25C42	25CB	38.84	25N	25C42	23CA	73.38
25N	25C42	24N	41.94	25N	25C42	26N	41.41
25N	25C42	25CA	19.88	25N	25C42	24C	16.17
25N	25C42	24CA	33.54	25N	25C42	26NE1	68.49
25N	25C42	25C	31.76	25N	25C42	26CG	65.77
25N	25C42	26CB	67.36	26CD1	25C42	65CA	59.61
26CD1	25C42	25CB	93.02	26CD1	25C42	23CA	96.28
26CD1	25C42	24N	69.60	26CD1	25C42	26N	46.05
26CD1	25C42	25CA	73.18	26CD1	25C42	24C	52.84
26CD1	25C42	24CA	52.35	26CD1	25C42	65N	66.14
26CD1	25C42	26NE1	15.58	26CD1	25C42	25C	60.30
26CD1	25C42	26CG	12.37	26CD1	25C42	5 <b>6N</b>	42.73
26CD1	25C42	65C	47.92	26CD1	25C42	26CB	29.69
65CA	25C42	23CA	85.42	65CA	25C42	24N	91.42

TABLE XVI								
65CA	25C42	24CA	88.16	65CA	25C42	65N	17.82	
65CA	25C42	26NE1	51.70	65CA	25C42	26CG	67.28	
65CA	25C42	6 <b>6N</b>	29.94	65CA	25C42	65C	14.64	
65CA	25C42	26CB	78.77	25CB	25C42	23CA	97.37	
25CB	25C42	24N	76.40	25CB	25C42	26N	48.95	
25CB	25C42	25CA	21.77	25CB	25C42	24C	54.30	
25CB	25C42	24CA	72.26	25CB	25C42	25C	33.23	
25CB	25C42	26CG	86.02	25CB	25C42	26CB	76.81	
23CA	25C42	24N	35.27	23CA	25C42	25CA	90.57	
23CA	25C42	24C	69.58	23CA	25C42	24CA	53.60	
23CA	25C42	65N	67.73	23CA	25C42	26NE1	82.24	
23CA	25C42	65C	95.61	24N	25C42	26N	75.74	
24N	25C42	25CA	61.52	24N	25C42	24C	34.63	
24N	25C42	24CA	19.12	24N	25C42	65N	78.18	
24N	25C42	26NE1	60.00	24N	25C42	25C	71.96	
24N	25C42	26CG	77.77	24N	25C42	65C	93.70	
24N	25C42	26CB	89.98	26N	25C42	25CA	34.07	
26N	25C42	24C	41.33	26N	25C42	24CA	58.54	
26N	25C42	26NE1	58.48	26N	25C42	25C	16.06	
26N	25C42	26CG	37.28	26N	25C42	66N	80.08	
26N	25C42	65C	91.09	26N	25C42	26CB	29.32	
25CA	25C42	24C	33.53	25CA	25C42	24CA	52.86	
25CA	25C42	26NE1	80.09	25CA	25C42	25C	18.89	
25CA	25C42	26CG	68.25	25CA	25C42	26CB	63.36	
24C	25C42	24CA	20.14	24C	25C42	65N	97.32	
24C	25C42	26NE1	52.89	24C	25C42	25C	38.23	
24C -	25C42	26CG	54.87	24C	25C42	66N	95.24	
24C	25C42	65C	96.26	24C	25C42	26CB	60.96	
24CA	25C42	65N	79.58	24CA	25C42	26NE1	45.69	
24CA	25C42	25C	58.08	24CA	25C42	26CG	59.27	
24CA	25C42	66N	89.86	24CA	25C42	65C	85.80	
24CA	25C42	26CB	70.86	65N	25C42	26NE1	54.04	
65N	25C42	26CG	76.36	65N	25C42	66N	46.51	
65N	25C42	65C	30.57	65N	25C42	26CB	90.76	
26NE1	25C42	25C	70.82	26NE1	25C42	26CG	27.95	
26NE1	25C42	66N	44.28	26NE1	25C42	65C	43.42	
26NE1	25C42	26CB	45.27	25C	25C42	26CG	52.79	
25C	25C42	66N	96.14	25C	25C42	26CB	45.17	

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	TABLE XVI								
26CG	25C42	66N	44.88	26CG	25C42	65C	53.98		
26CG	25C42	26CB	17.32	66N	25C42	65C	15.94		
66N	25C42	26CB	51.51	65C	25C42	26CB	64.35		
66N	25N43	65CA	39.20	6 <b>6N</b>	25 <b>N4</b> 3	660	42.83		
66N	25 <b>N4</b> 3	65C	19.66	66N	25N43	66C	33.44		
66N	25N43	66CA	16.52	6 <b>6N</b>	25N43	640	67.91		
66N	25N43	650	24.48	6 <b>6N</b>	25N43	65N	47.82		
65CA	25N43	660	79.74	65CA	25N43	65C	22.72		
65CA	25N43	66C	72.52	65CA	25N43	66CA	54.97		
65CA	25N43	640	35.80	65CA	25 <b>N4</b> 3	650	27.77		
65CA	25 <b>N4</b> 3	65N	9.59	660	25N43	65C	62.50		
660	25N43	66°C	14.75	660	25N43	66CA	32.80		
660	25N43	650	66.04	660	25N43	65N	89.14		
65C	25N43	66C	52.38	65C	25N43	66CA	33.37		
65C	25 <b>N4</b> 3	640	48.27	65C	25N43	650	9.39		
65C	25N43	65N	29.76	66C	25 <b>N4</b> 3	66CA	19.82		
66C	25 <b>N4</b> 3	640	99.32	66C	25N43	650	54.13		
66C	25 <b>N4</b> 3	65N	81.27	66CA	25N43	640	79.51		
66CA	25 <b>N4</b> 3	650	34.33	66CA	25N43	65N	62.98		
640	25 <b>N4</b> 3	650	45.20	640	25N43	65N	26.86		
650	25N43	65N	32.42						

TABLE XVII

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-

[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
67CE2	25C1	67CD2	21.66	67CE2	25C1	60ND2	88.40
67CE2	25C1	67CZ	15.07	67CE2	25C1	2310H2	72.49
67CE2	25C1	670H	28.98	67CE2	25C1	66CA	80.69
67CE2	25C1	67CG	26.79	67CE2	25C1	70ND2	78.17
67CD2	25C1	60ND2	67.38	67CD2	25C1	67CZ	33.45
67CD2	25C1	2310H2	62.87	67CD2	25C1	670H	49.60
67CD2	25C1	66CA	68.97	67CD2	25C1	67CG	10.46
67CD2	25C1	70ND2	58.29	60ND2	25C1	67CZ	95.36
60ND2	25C1	590	49.64	60ND2	25C1	2310H2	73.76
60ND2	25C1	66CA	42.03	60ND2	25C1	650	60.94
60ND2	25C1	67CG	61.96	60ND2	25C1	70ND2	36.62
67CZ	25C1	2310H2	87.30	67CZ	25C1	670H	16.70
67CZ	25C1	66CA	77.40	67CZ	25C1	67CG	34.47
67CZ	25C1	70ND2	91.66	590	25C1	2310H2	59.15
590	25C1	66CA	86.93	590	25C1	650	85.11
590	25C1	70ND2	43.49	2310H2	25C1	670H	96.03
2310H2	25C1	67CG	71.00	2310H2	25C1	70ND2	37.14
670H	25C1	66CA	87.65	670H	25C1	67CG	51.17
66CA	25C1	650	33.94	66CA	25C1	67CG	58.73
66CA	25C1	70ND2	75.50	650	25C1	67CG	92.09
650	25C1	70ND2	97.34	67CG	25C1	70ND2	60.11
67CE2	25C2	67CZ	20.70	67CE2	25C2	67CD2	20.73
67CE2	25C2	670H	36.29	57CE2	25C2	66CA	91.85
67CE2	25C2	67CE1	28.15	67CE2	25C2	60ND2	82.91
67CE2	25C2	6 <b>6</b> C	74.08	67CE2	25C2	67CG	28.19
67CE2	25C2	67N	€3.54	€7CZ	25C2	67CD2	36.53
67CZ	25C2	670H	19.80	67 <b>CZ</b>	25C2	66CA	91.39
67CZ	25C2	66N	97.52	67CZ	25C2	67CE1	14.08
67CZ	25C2	60ND2	96.48	3 <b>7CZ</b>	25C2	6 <b>6</b> C	73.78
67CZ	25C2	67CG	36.03	67 <b>CZ</b>	25C2	67N	68.45

	TABLE XVII								
67CD2	25C2	670H	55.27	67CD2	25C2	66CA	75.10		
67CD2	25C2	66N	89.93	67CD2	25C2	67CE1	35.97		
67CD2	25C2	60ND2	62.30	67CD2	25C2	66C	58.50		
67CD2	25C2	67CG	13.99	67CD2	25C2	67N	45.40		
670H	25C2	67CE1	30.01	670H	25C2	66C	90.02		
670H	25C2	67CG	55.88	670H	25C2	67N	87.15		
66CA	25C2	650	39.05	66CA	25C2	66N	18.31		
66CA	25C2	65C	31.60	66CA	25C2	67CE1	77.85		
66CA	25C2	60ND2	42.40	66CA	25C2	66C	17.99		
66CA	25C2	67CG	63.85	66CA	25C2	67N	29.89		
650	25C2	66N	29.44	650	25C2	65C	15.02		
650	25C2	60ND2	62.06	650	25C2	66C	56.82		
650	25C2	67N	68.16	6 <b>6N</b>	25C2	65C	16.65		
66N	25C2	67CE1	83.45	6 <b>6N</b>	25C2	60ND2	59.05		
66N	25C2	66C	31.58	66N	25C2	67CG	77.19		
66N	25C2	67N	46.44	65C	25C2	67CE1	99.01		
65C	25C2	60ND2	65.42	65C	25C2	66C	47.55		
65C	25C2	67CG	93.66	65C	25C2	67N	61.39		
67CE1	25C2	60ND2	88.93	67CE1	25C2	66C	60.58		
67CE1	25C2	67CG	29.84	67CE1	25C2	67N	57.26		
60ND2	25C2	66C	44.34	60ND2	25C2	67CG	60.48		
60ND2	25C2	67N	35.75	66C	25C2	67CG	46.38		
66C	25C2	67N	15.93	67CG	25C2	67N	35.42		
650	25C3	65C	17.88	650	25C3	66CA	41.69		
650	25C3	66N	32.43	650	25C3	61CB	43.31		
650	25C3	65CA	28.80	650	25C3	61CG	61.48		
650	25C3	610D1	68.40	65C	25C3	66CA	35.13		
65C	25C3	.66N	18.49	65C	25C3	61CB	59.01		
65C	25C3	65CA	16.90	65C	25C3	61CG	77.19		
65C	25C3	610D1	85.63	66CA	25C3	66N	20.18		
66CA	25C3	67CE2	75.24	6 <b>6CA</b>	25C3	61CB	82.37		
66CA	25C3	65CA	50.05	66CA	25C3	61CG	99.19		
66CA	25C3	67CZ	76.99	66CA	25C3	670H	90.94		
6EN	25C3	67CE2	90.18	6 <b>6N</b>	25C3	61CB	75.69		
66N	25C3	65CA	30.59	6 <b>6N</b>	25C3	61CG	93.89		
66N	25C3	67CZ	87.56	6 <b>6N</b>	25C3	670H	98.55		
67CE2	25C3	67CZ	16.0ó	67CE2	25C3	670H	28.37		
61CB	25C3	65CA	59.68	61CB	25C3	61CG	18.24		

			TABI	LE XVII			
61CB	25C3	610D1	28.38	65CA	25C3	61CG	76.44
65CA	25C3	610D1	88.03	61CG	25C3	610D1	14.65
67CZ	25C3	670H	15.78	650	25C4	61CG	81.72
650	25C4	61CB	56.74	650	25C4	610D1	92.04
650	25C4	610D2	90.06	650	25C4	65C	13.06
650	25C4	61N	47.80	650	25C4	61CA	52.82
650	25C4	66CA	34.06	650	25C4	66N	23.80
650	25C4	60C	59.87	650	25C4	65CA	23.94
61CG	25C4	61CB	24.99	61CG	25C4	610D1	20.63
61CG	25C4	610D2	17.88	61CG	25C4	65C	91.97
61CG	25C4	61N	50.91	61CG	25C4	61CA	33.78
61CG	25C4	60C	56.14	61CG	25C4	65CA	87.08
61CB	25C4	610D1	39.37	61CB	25C4	610D2	36.30
61CB	25C4	65C	67.18	61CB	25C4	61N	35.38
61CB	25 <b>C4</b>	61CA	16.65	61CB	25C4	66CA	87.99
61CB	25C4	66N	80.49	61CB	25C4	60C	47.22
61CB	25C4	65CA	64.13	610D1	25C4	610D2	33.84
610D1	25C4	61N	49.82	610D1	25C4	61CA	39.25
610D1	25C4	60C	48.15	610D2	25C4	65C	97.5 <del>9</del>
610D2	25C4	61N	67.70	610D2	25C4	61CA	49.40
610D2	25C4	60C	73.98	610D2	25C4	65CA	88.32
65C	25C4	61N	60.83	65C	25C4	61CA	65.24
65C	25C4	66CA	30.77	65C	25C4	66N	15.04
65C	25C4	60C	72.43	65C	25C4	65CA	16.09
61N	25C4	61CA	19.04	61N	25C4	66CA	65.40
61N	25C4	66N	67.36	61N	25C4	60C	15.56
61N	25C4	65CA	68.40	61CA	25C4	66CA	78.91
61CA	25C4	66N	75.90	61CA	25C4	60C	30.75
61CA	25C4	65CA	67.27	66CA	25C4	66N	17.65
66CA	25C4	60C	69.36	66CA	25C4	65CA	45.82
66 <b>N</b>	25C4	60C	75.84	66N	25C4	65CA	28.52
60C	25C4	65CA	82.21	61CD1	25 <b>C</b> 5	61CG	21.34
610D1	25C5	61CB	38.98	610D1	25C5	61N	54.59
610D1	25C5	650	86.61	61001	25C5	60C	57.24
610D1	25C5	61002	31.11	61001	25C5	590	91.41
610D1	25C5	60CA	77.36	610D1	25C5	61CA	39.29
610D1	25 <b>C</b> 5	600	47.14	61CG	25C5	61CB	23.42
61CG	25C5	61N	53.21	61CC	25C5	650	71.11

TABLE XVII								
61CG	25C5	60C	63.55	61CG	25C5	610D2	15.46	
61CG	25C5	60CA	84.04	61CG	25C5	61CA	34.30	
61CG	25C5	600	58.88	61CB	25C5	61N	36.38	
61CB	25C5	650	48.37	61CB	25C5	60C	52.26	
61CB	25C5	610D2	34.12	61CB	25C5	60CA	70.23	
61CB	25C5	61CA	19.16	61CB	25C5	600	54.58	
61CB	25C5	60ND2	92.51	61N	25C5	. 650	46.30	
61N	25C5	60C	18.83	61N	25C5	610D2	67.79	
61N	25C5	590	68.18	61N	25C5	60CA	33.92	
61N	25C5	61CA	19.02	61N	25C5	600	29.73	
61N	25C5	60ND2	59.73	650	25C5	60C	62.13	
650	25C5	610D2	76.92	650	25C5	590	97.98	
650	25C5	60CA	63.97	650	25C5	61CA	50.38	
650	25C5	600	75.74	650	25C5	60ND2	60.02	
60C	25C5	610D2	79.00	60C	25C5	590	50.32	
60C	25C5	60CA	20.70	60C	25C5	61CA	33.16	
60C	25C5	600	15.46	60C	25C5	60ND2	54.15	
610D2	25C5	60CA	99.43	610D2	25C5	61CA	48.77	
610D2	25C5	600	73.92	590	25C5	60CA	36.41	
590	25C5	61CA	83.15	590	25C5	600	49.36	
590	25C5	60ND2	45.49	60CA	25C5	61CA	51.88	
60CA	25C5	600	31.33	60CA	25C5	60ND2	34.90	
61CA	25C5	600	36.38	61CA	25C5	60ND2	78.49	
600	25C5	60ND2	66.22	590	25C6	610D1	84.33	
590	25C6	60CA	37.74	590	25C6	60ND2	52.70	
590	25C6	60C	49.43	590	25C6	59C	7.16	
590	25C6	2310H2	61.67	590	25C6	650	96.01	
590	25C6	61N	65.44	590	25C6	61CG	94.50	
610D1	25C6	60CA	67.00	610D1	25C6	60C	47.52	
610D1	25C6	59C	79.80	610D1	25C6	650	65.95	
610D1	25C6	61N	43.14	610D1	25C6	61CG	14.44	
60CA	25C6	60ND2	38.07	50CA	25C6	60C	19.60	
60CA	25 <b>C</b> 6	59C	30.68	60CA	25C6	2310H2	90.52	
60CA	25C6	650	58.49	60CA	25C6	61N	31.11	
60CA	25C6	61CG	70.15	60ND2	25C6	60C	55.78	
60ND2	25C6	59C	49.33	60ND2	25C6	2310H2	69.41	
60MD2	25C6		60.19		25C6		59.59	
60ND2	25C6	67CE2	71.07	60C	25C6	59C	42.53	

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TABLE XVII							
60C	25C6	650	53.64	60C	25C6	61N	16.42
60C	25C6	61CG	50.89	59C	25C6	2310H2	67.54
59C	25C6	650	89.09	59C	25C6	61N	58.40
59C	25C6	61CG	89.02	2310H2	25C6	67CE2	58.91
650	25C6	61N	38.82	650	25C6	61CG	54.53
650	25C6	67CE2	90.16	61N	25C6	61CG	41.32
650	25C7	65C	20.61	650	25C7	6 <b>6N</b>	35.57
650	25C7	65CA	37.59	650	25C7	640	63.61
650	25C7	66CA	40.23	650	25C7	65N	36.78
650	25C7	64C	50.16	65C	25C7	66N	20.13
65C	25C7	65CA	22.56	65C	25C7	640	58.37
65C	25C7	66CA	34.64	65C	25C7	65N	30.89
65C	25C7	64C	46.81	66N	25C7	65CA	35.80
66N	25C7	640	74.14	66N	25C7	66CA	19.87
66N	25C7	65N	48.83	66N	25C7	64C	64.44
65CA	25C7	640	38.35	65CA	25C7	66CA	54.61
65CA	25C7	65N	16.17	65CA	25C7	64C	29.70
640	25C7	66CA	92.52	640	25C7	65N	28.03
640	25C7	64C	13.53	66CA	25C7	65N	65.51
66CA	25C7	64C	81.44	66CA	25C7	670H	89.07
65N	25C7	64C	15.93	66N	2508	65C	20.16
66N	2508	66CA	21.24	6 <b>6N</b>	2508	650	33.23
66N	2508	65CA	33.75	66N	2508	67CE1	98.44
6 <b>6</b> N	2508	67CE2	98.98	66N	2508	66C	34.36
66N	2508	660	36.02	6 <b>6N</b>	2508	640	65.86
65C	2508	66CA	35.77	65C	2508	650	17.66
65C	2508	65CA	20.48	65C	2508	6 <b>6</b> C	52.97
65C	2508	660	56.18	65C	2508	640	48.74
66CA	2508	650	39.76	66CA	2508	67CZ	88.60
66CA	2508	65CA	53.75	65CA	2508	67CE1	82.16
66CA	2508	67CE2	78.01	66CA	2508	66C	19.45
66CA	2508	660	29.94	65CA	2508	640	84.46
670H	2508	67CZ	19.19	670H	2508	67CE1	31.81
670H	2508	67CE2	31.82	570H	2508	66C	93.01
670H	2508	660	95.44	650	2508	65CA	33.40
650	2508	66C	59.16	650	2508	660	66.79
650	2508	540	51.69	67CZ	2508	67CE1	17.96
67CZ	2508	67CE2	17.62	67CZ	2508	66C	73.92

			TA	BLE XVII			
67CZ	2508	660	77.47	65CA	2508	66C	67.93
65CA	2508	660	66.32	65CA	2508	640	32.46
67CE1	2508	67CE2	30.82	67CE1	2508	66C	64.46
67CE1	2508	660	64.18	67CE2	2508	66C	67.34
67CE2	2508	660	75.18	66C	2508	660	15.05
660	2508	640	98.36	66N	25C9	65C	19.56
66N	25C9	65CA	35.74	66N	25C9	66CA	18.50
66N	25C9	650	29.44	66N	25C9	660	37.27
66N	25C9	640	68.91	66N	25C9	66C	32.34
66N	25C9	67CE1	91.45	66N	25C9	67CZ	94.14
65C	25 <b>C9</b>	65CA	21.89	65C	25C9	66CA	33.21
65C	25C9	650	14.71	65C	25C9	660	56.83
65C	25C9	640	50.85	65C	25C9	66C	50.46
65CA	25C9	66CA	53.09	65CA	25C9	650	31.99
65CA	25C9	660	69.79	65CA	25 <b>C</b> 9	640	34.59
65CA	25C9	66C	67.96	66CA	25C9	650	35.79
66CA	25C9	660	30.43	66CA	25C9	640	83.99
66CA	25C9	66C	18.68	66CA	25 <b>C</b> 9	670H	90.57
66CA	25C9	67CE1	74.02	66CA	25C9	67CZ	75.67
650	25C9	660	64.42	650	25C9	640	51.51
650	25C9	66C	54.47	650	25C9	67CZ	97.86
660	25C9	66C	15.18	660	25C9	670H	88.11
660	25C9	67CE1	62.23	660	25C9	67CZ	72.01
66C	25C9	670H	81.72	66C	25C9	67CE1	59.79
66C	25C9	67CZ	65.61	670H	25C9	67CE1	28.34
670H	25C9	67CZ	16.24	67CE1	25C9	67CZ	16.27
65CA	25010	640	37.93	65CA	25010	65C	19.98
65CA	25010	6 <b>6N</b>	32.25	640	25010	65C	52.12
640	25010	6 <b>6N</b>	68.39	65C	25010	66N	17.19
66N	25N11	660	46.02	6 <b>6N</b>	25N11	66C	38.85
66N	25N11	66CA	20.73	EEN	25N11	65C	17.46
66N	25N11	65CA	33.59	8 <b>6N</b>	25N11	67CD1	91.49
66N	25N11	650	24.15	66.01	25N11	67N	48.38
660	25N11	66C	17.87	660	25N11	66CA	35.52
660	25N11	65C	63.48	860	25N11	67CE1	75.71
660	25N11	65CA	76.62	660	25N11	67CZ	83.40
660	25N11	67CD1	58.50	660	25N11	670H	99.66
660	25N11	650	67.57	660	25N11	67N	24.87

TABLE XVII									
66C	25N11	66CA	21.72	66C	25N11	65C	54.97		
66C	25N11	67CE1	70.15	66C	25N11	65CA	72.37		
66C	25N11	67CZ	73.23	66C	25N11	67CD1	54.55		
66C	25N11	670H	88.69	66C	25N11	650	55.57		
6 <b>6</b> C	25N11	67N	11.56	66CA	25N11	65C	34.29		
66CA	25N11	67CE1	83.75	66CA	25N11	65CA	53.45		
66CA	25N11	67CZ	81.15	66CA	25N11	67CD1	71.00		
66CA	25N11	670H	93.79	66CA	25N11	650	33.86		
66CA	25N11	67N	28.86	65C	25N11	65CA	20.50		
65C	25N11	650	12.17	65C	25N11	67N	63.15		
67CE1	25 <b>N1</b> 1	67CZ	16.97	67CE1	25N11	67CD1	17.33		
67CE1	25N11	670H	28.98	67CE1	25N11	67N	58.73		
65CA	25N11	650	29.33	65CA	25N11	67N	81.90		
67CZ	25N11	67CD1	29.85	67CZ	25N11	670H	16.29		
67CZ	25N11	650	95.29	67CZ	25N11	67N	61.81		
67CD1	25 <b>N11</b>	670H	44.88	67CD1	25N11	650	98.49		
67CD1	25N11	67N	43.62	670H	25N11	67N	77.48		
650	25N11	67N	61.50	660	25C12	66N	39.81		
660	25C12	66C	13.28	660	25C12	65CA	68.49		
660	25C12	66CA	28.41	660	25C12	65C	53.90		
66N	25C12	66C	32.53	6 <b>6N</b>	25C12	65CA	30.03		
66N	25C12	66CA	16.05	6 <b>6N</b>	25C12	65C	14.10		
1610	25C12	161C	14.53	66C	25C12	65CA	62.55		
66C	25C12	66CA	17.90	66C	25C12	65C	45.95		
65CA	25C12	66CA	45.67	65CA	25C12	65C	17.89		
66CA	25C12	65C	28.39	660	25C13	6 <b>6N</b>	46.61		
660	25C13	65CA	81.12	660	25C13	26CB	42.40		
660	25C13	66C	12.53	660	25C13	26CD1	59.74		
660	25C13	65C	61.86	660	25C13	66CA	31.09		
660	25C13	26CG	46.38	6 <b>6</b> 0	25C13	2 <b>6N</b>	73.59		
6 <b>6N</b>	25C13	65CA	34.83	66N	25C13	26CB	65.68		
6 <b>6N</b>	25C13	66C	35.92	6 <b>6N</b>	25C13	26CD1	50.75		
66.N	25C13	65C	15.26	6 <b>6N</b>	25C13	66CA	16.54		
66N	25C13	26CG	52.33	6 <b>6N</b>	25C13	26N	87.54		
25SG	25C13	26CB	74.78	25 <i>S</i> G	25C13	26CD1	73.77		
25 <i>S</i> G	25C13	26CG	78.01	25SG	25C13	26N	43.94		
25SG	25C13	1610	62.15	65CA	25C13	26CB	87.36		
65CA	25C13	66C	70.75	65CA	25C13	26CD1	57.53		

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TABLE XVII								
65CA	25C13	65C	20.04	65CA	25C13	66CA	51.26	
65CA	25C13	26CG	69.09	65CA	25C13	26N	94.25	
26CB	25C13	66C	51.08	26CB	25C13	26CD1	34.05	
26CB	25C13	65C	76.87	26CB	25C13	66CA	59.32	
26CB	25C13	26CG	18.75	26CB	25C13	26N	31.21	
66C	25C13	26CD1	60.72	66C	25C13	65C	50.94	
66C	25C13	66CA	19.59	66C	25C13	26CG	50.32	
66C	25C13	26N	81.92	26CD1	25C13	65C	54.60	
26CD1	25C13	66CA	55.37	26CD1	25C13	26CG	16.44	
26CD1	25C13	26N	38.73	65C	25C13	66CA	31.37	
65C	25C13	26CG	60.91	65C	25C13	26N	93.30	
66CA	25C13	26CG	51.05	66CA	25C13	26N	86.68	
26CG	25C13	26N	35.89	25 <b>S</b> G	25N14	230	84.00	
25SG	25N14	26CD1	79.14	25SG	25N14	1610	68.92	
25SG	25 <b>N14</b>	26N	43.55	25SG	25N14	23C	74.81	
25SG	25N14	26CB	70.46	25 <b>S</b> G	25 <b>N14</b>	26CG	77.79	
65CA	25N14	6 <b>6N</b>	36.05	65CA	25 <b>N14</b>	230	53.08	
65CA	25N14	26CD1	61.64	65CA	25N14	65C	18.94	
65CA	25N14	660	73.48	65CA	25N14	65N	12.48	
65CA	25N14	26N	99.87	65CA	25 <b>N14</b>	23C	64.69	
65CA	25N14	26CB	84.32	65CA	25N14	26CG	69.23	
66N	25N14	230	73.08	66N	25N14	26CD1	49.53	
6 <b>6N</b>	25 <b>N14</b>	65C	17.28	66N	25N14	660	37.50	
66N	25N14	65N	47.78	66N	25 <b>N14</b>	26N	82.41	
66N	25N14	23C	86.05	6 <b>6N</b>	25N14	26CB	56.95	
66N	25N14	26CG	48.25	230	25N14	26CD1	46.64	
230	25N14	65C	63.98	230	25N14	660	96.33	
230	25N14	65N	44.52	230	25N14	26N	67.73	
230	25N14	23C	13.02	230	25N14	26CB	76.26	
230	25N14	26CG	60.45	26CD1	25 <b>N14</b>	65C	55.15	
26CD1	25 <b>N14</b>	660	52.89	26CD1	25N14	65N	64.11	
26CD1	25 <b>N14</b>	26N	38.34	26CD1	25N14	23C	54.90	
26CD1	25 <b>N14</b>	26CB	30.70	26CD1	25 <b>N14</b>	26CG	13.86	
65C	25 <b>N14</b>	660	54.77	65C	25N14	65N	31.08	
65C	25N14	26N	92.32	6 <b>5</b> C	25N14	23C	76.79	
65C	25N14	26CB	70.71	65¢	25N14	26CG	58.49	
660	25N14	6 <b>5N</b>	84.80	660	25N1.4	26N	64.25	
660	25N14	26CB	35.42	660	25N14	26CG	41.33	

			TA	BLE XVII			
65N	25N14	23C	54.87	65N	25N14	26CB	90.63
65N	25N14	26CG	74.10	26N	25N14	23C	67.48
26N	25N14	26CB	29.19	26N	25N14	26CG	34.32
23C	25N14	26CB	82.14	23C	25N14	26CG	68.17
26CB	25N14	26CG	17.47	1610	25C15	25 <b>S</b> G	89.42
1610	25C15	161C	13.49	1610	25C15	162CA	34.91
1610	25C15	162N	23.43	1610	25C15	161CA	23.63
25SG	25C15	161C	88.53	25SG	25C15	162CA	57.17
25SG	25C15	162N	74.53	161C	25C15	162CA	31.37
161C	25C15	162N	14.79	161C	25C15	161CA	16.28
162CA	25C15	162N	17.76	162CA	25C15	161CA	46.94
162N	25C15	161CA	29.37	65CA	25C15	2300H2	72.79
25 <b>S</b> G	25C16	26N	61.11	25SG	25C16	25N	59.97
25 <i>S</i> G	25C16	25CB	23.02	25SG	25C16	25CA	42.42
25 <i>S</i> G	25C16	26CB	88.58	25SG	25C16	24N	92.96
25SG	25C16	25C	48.34	25 <i>S</i> G	25C16	24C	74.43
25SG	25C16	26CA	71.25	25 <i>S</i> G	25C16	24CA	91.58
230	25C16	26CD1	61.77	230	25C16	26N	94.53
230	25C16	2 <b>5N</b>	67.24	230	25C16	23C	18.20
230	25C16	65CA	58.52	230	25C16	25CA	87.06
230	25C16	26CG	77.13	230	25C16	26NE1	47.29
230	25C16	26CB	97.08	230	25C16	6 <b>6N</b>	79.63
230	25C16	24N	30.53	230	25C16	23CA	32.27
230	25C16	25C	94.82	230	25C16	24C	56.19
230	25C16	26CA	99.54	230	25C16	24CA	36.83
230	25C16	65C	67.39	230	25C16	65N	45.21
26CD1	25C16	26N	51.13	26CD1	25C16	25N	68.54
26CD1	25C16	23C	73.51	26CD1	25C16	65CA	68.04
26CD1	25C16	25CB	97.25	26CD1	25C16	25CA	76.12
26CD1	25C16	26CG	15.44	26CD1	25C16	26NE1	15.45
26CD1	25C16	26CB	35.76	26CD1	25C16	66N	52.08
26CD1	25C16	24N	69.01	26CD1	25C16	23CA	92.69
26CD1	25C16	25C	63.96	26CD1	25C16	24C	55.05
26CD1	25C16	26CA	44.91	26CD1	25C16	24CA	52.83
26CD1	25C16	65C	56.87	26CD1	25C16	660	54.10
26CD1	25C16	65N	69.48	26N	25C16	25N	44.67
26N	25C16	23C	93.12	26N	25C16	25CB	51.34
26N	25C16	25CA	35.79	26N	25C16	26CG	42.62

TABLE XVII								
26N	25C16	26NE1	63.79	26N	25C16	26CB	34.80	
26N	25C16	66N	93.89	26N	25C16	24N	77.38	
26N	25C16	25C	16.11	26N	25C16	24C	44.64	
26N	25C16	26CA	16.33	26N	25C16	24CA	61.13	
26N	25C16	660	70.60	25N	25C16	23C	56.44	
25N	25C16	25CB	37.16	25N	25C16	25CA	19.86	
25N	25C16	26CG	71.75	25N	25C16	26NE1	70.29	
25N	25C16	26CB	75.60	25N	25C16	24N	39.16	
25N	25C16	23CA	68.19	25N	25C16	25C	33.83	
25N	25C16	24C	15.28	25N	25C16	26CA	60.18	
25N	25C16	24CA	31.81	23C	25C16	65CA	74.98	
23C	25C16	25CB	86.20	23C	25C16	25CA	75.86	
23C	25C16	26CG	87.97	23C	25C16	26NE1	60.92	
23C	25C16	66N	97.84	23C	25C16	24N	17.30	
23C	25C16	23CA	19.93	23C	25C16	25C	88.48	
23C	25C16	24C	49.42	23C	25C16	24CA	32.05	
23C	25C16	65C	85.33	23C	25C16	65N	60.76	
65CA	25C16	26CG	76.49	65CA	25C16	26NE1	57.92	
65CA	25C16	26CB	90.45	65CA	25C16	6 <b>6N</b>	33.37	
65CA	25C16	24N	89.01	65CA	25C16	23CA	76.80	
65CA	25C16	24CA	90.04	65CA	25C16	65C	17.45	
65CA	25C16	660	68.34	65 <b>CA</b>	25C16	65N	14.75	
25CB	25C16	25CA	21.65	25CB	25C16	26CG	93.09	
25CB	25C16	26CB	85.06	25CB	25C16	24N	70.81	
25CB	25C16	23CA	89.00	25CB	25C16	25C	35.50	
25CB	25C16	24C	52.00	25 <b>CB</b>	25C16	26CA	65.90	
25CB	25C16	24CA	68.59	25 <b>CA</b>	25C16	26CG	73.95	
25CA	25C16	26NE1	82.49	25 <b>CA</b>	25C16	26CB	70.49	
25CA	25C16	24N	58.71	2 <b>5CA</b>	25C16	23CA	85.38	
25CA	25C16	25C	20.38	2 <b>5CA</b>	25C16	24C	32.49	
25CA	25C16	26CA	52.03	25CA	25C16	24CA	51.06	
26CG	25C16	25NE1	30.69	26 <b>CG</b>	25C16	26CB	20.39	
26CG	25C16	66N	52.82	26CG	25C16	24N	81.30	
26CG	25C16	25C	57.74	26CG	25C16	24C	60.85	
26CG	25C16	26CA	32.36	26CG	25C16	24CA	63.73	
26CG	25C16	65C	62.21	26 <b>CG</b>	25C16	660	43.58	
26CG	25C16	65N	80.99	26NE1	25C16	26CB	51.09	
26NE1	25C16	66N	51.16	26NE1	25C16	24N	59.98	

			TAE	BLE XVII			
26NE1	25C16	23CA	79.09	26NE1	25C16	25C	74.44
26NE1	25C16	24C	55.27	26NE1	25C16	26CA	59.77
26NE1	25C16	24CA	46.88	26NE1	25C16	65C	50.71
26NE1	25C16	660	63.77	26NE1	25C16	65N	56.44
26CB	25C16	66N	60.55	26CB	25C16	24N	96.31
26CB	25C16	25C	50.81	26CB	25C16	24C	69.20
26CB	25C16	26CA	19.18	26CB	25C16	24CA	77.92
26CB	25C16	65C	73.91	26CB	25C16	660	36.73
26CB	25C16	65N	98.03	66N	25C16	26CA	79.58
66N	25C16	24CA	97.14	66N	25C16	65C	16.15
66N	25C16	660	35.16	66N	25C16	65N	46.29
24N	25C16	23CA	32.77	24N	25C16	25C	71.39
24N	25C16	24C	32.88	24N	25C16	26CA	88.94
24N	25C16	24CA	18.41	24N	25C16	65C	96.24
24N	25C16	65 <b>N</b>	75.67	23CA	25C16	24C	65.31
23CA	25C16	24CA	50.41	23CA	25C16	65C	91.28
23CA	25C16	65N	62.12	25C	25C16	24C	39.37
25C	25C16	26CA	31.92	25C	25C16	24CA	58.36
25C	25C16	660	85.95	24C	25C16	26CA	57.54
24C	25C16	24CA	19.36	24C	25C16	65 <b>N</b>	97.77
26CA	25C16	24CA	71.17	26CA	25C16	65C	92.26
26CA	25C16	660	54.28	24CA	25C16	65C	91.93
24CA	25C16	65N	79.37	65C	25C16	660	51.31
65C	25C16	65N	30.19	660	25C16	65N	81.41
25SG	25C17	25CB	33.36	25 <b>S</b> G	25C17	25 <b>N</b>	69.53
25SG	25C17	25CA	47.13	25 <i>S</i> G	25C17	26N	51.70
25SG	25C17	25C	42.10	25 <i>S</i> G	25C17	1610	81.99
25SG	25C17	162ND1	45.09	25 <b>S</b> G	25C17	24C	78.33
25SG	25C17	26CD1	91.04	25 <b>SG</b>	25C17	19NE2	91.40
25SG	25C17	24CA	96.69	25 <b>S</b> G	25C17	163N	23.29
25SG	25C17	162CA	48.75	25CB	25C17	25 <b>N</b>	43.65
25CB	25C17	25CA	23.36	25CB	25C17	23C	97.56
25CB	25C17	26N	52.35	25CB	25C17	24N	79.60
25CB	25C17	25C	34.92	25CB	25C17	162ND1	47.80
25CB	25C17	24C	54.50	25CB	25C17	26CD1	89.27
25CB	25C17	19NE2	59.46	25CB	25C17	24CA	71.68
25CB	25C17	163N	55.20	25CB	25C17	162CA	73.99
25N	25C17	25CA	22.58	25N	25C17	230	63.36

		ТД	BLE XVII		•
25N	25C17 23C	56.78	25N	25C17 26N	42.10
25N	25C17 23CA	72.74	25N	25C17 24N	39.75
25N	25C17 25C	33.56	25N	25C17 162ND1	89.46
25N	25C17 24C	11.07	25N	25C17 26CD1	57.00
25N	25C17 19NE2	52.78	25 <b>N</b>	25C17 24CA	28.26
25N	25C17 163N	92.62	25CA	25C17 230	85.24
25CA	25C17 23C	79.30	25CA	25C17 26N	36.07
25CA	25C17 23CA	93.96	25CA	25C17 24N	61.96
25CA	25C17 25C	19.74	25CA	25C17 162ND1	71.14
25CA	25C17 24C	32.47	25CA	25C17 26CD1	67.46
25CA	25C17 19NE2	60.34	25CA	25C17 24CA	50.61
25CA	25C17 163N	70.36	25CA	25C17 162CA	94.23
230	25C17 23C	18.25	230	25C17 26N	78.82
230	25C17 23CA	34.40	230	25C17 24N	31.56
230	25C17 25C	86.35	230	25C17 24C	52.82
230	25C17 26CD1	45.95	230	25C17 19NE2	76.58
230	25C17 230OH2	65.85	230	25C17 24CA	35.23
23C	25C17 26N	84.24	23C	25C17 23CA	21.57
23C	25C17 24N	17.96	23C	25C17 25C	86.44
23C	25C17 24C	48.18	23C	25C17 26CD1	59.52
23C	25C17 19NE2	58.61	23C	25С17 230ОН2	60.40
23C	25C17 24CA	30.09	26N	25C17 24N	73.12
26N	25C17 25C	17.65	26N	25C17 162ND1	93.89
26N	25C17 24C	42.62	26N	25C17 26CD1	40.31
26N	25C17 19NE2	93.61	26N	25C17 24CA	56.29
26N	25C17 163N	67.72	2 <b>6N</b>	25C17 162CA	96.10
23CA	25C17 24N	33.68	23CA	25C17 24C	66.27
23CA	25C17 26CD1	79.75	23CA	25C17 19NE2	53.84
23CA	25C17 230OH2	40.17	23CA	25C17 24CA	49.58
24N	25C17 25C	71.51	24N	25C17 24C	32.59
24N	25C17 26CD1	59.47	24N	25C17 19NE2	47.60
24N	25C17 230OH2	73.74	24N	25C17 24CA	16.96
25C	25C17 162ND1	79.14	25C	25C17 24C	38.92
25C	25C17 26CD1	<b>55.4</b> 5	25C	25C17 19NE2	79.80
25C	25C17 24CA	56.38	25C	25C17 163N	62.85
25C	25C17 162CA	90.50	1610	25C17 162ND1	62.06
1.610	25C17 230OH2	90.45	1610	25C17 163N	60.08
1610	25C17 162CA	33.46	162ND1	25C17 19NE2	71.55

			TA	BLE XVII			
162ND1	25C17	163N	44.58	162ND1	25C17	162CA	38.78
24C	25C17	26CD1	48.36	24C	25C17	19NE2	57.22
24C	25C17	24CA	18.38	26CD1	25C17	24CA	46.21
19NE2	25C17	2300H2	73.47	19NE2	25C17	24CA	57.12
2300H2	25C17	24CA	89.69	163N	25C17	162CA	28.44
25 <i>S</i> G	25C18	1610	91.87	25SG	25C18	25CB	20.14
25 <i>S</i> G	25C18	23CA	98.61	25SG	25C18	161C	85.38
25SG	25C18	230	83.74	25SG	25C18	23C	83.16
25 <i>S</i> G	25C18	162ND1	47.66	25 <b>S</b> G	25C18	162CA	57.49
25 <i>S</i> G	25C18	25N	40.60	1610	25C18	161C	8.04
1610	25C18	162ND1	67.18	1610	25C18	162CA	35.14
2300H2	25C18	23CA	43.13	2300H2	25C18	230	66.88
2300H2	25C18	23C	61.23	25CB	25C18	23CA	84.02
25CB	25C18	161C	98.23	25CB	25C18	230	77.59
25CB	25C18	23C	72.21	25CB	25C18	162ND1	43.49
25CB	25C18	162CA	68.17	25CB	25C18	25N	30.01
23CA	25C18	230	30.55	23CA	25C18	23C	19.20
23CA	25C18	25N	58.06	161C	25C18	162ND1	65.74
161C	25C18	162CA	30.20	230	25C18	23C	15.25
230	25C18	25N	47.60	23C	25C18	25N	43.17
162ND1	25C18	162CA	39.32	162ND1	25C18	25N	72.26
162CA	25C18	25N	96.88	660	25C19	67CE1	68.17
660	25C19	67CD1	54.92	1600	25C19	161C	52.84
1600	25C19	161CA	35.03	1600	25C19	1610	63.95
1600	25C19	162N	57.36	67CE1	25C19	67CD1	17.17
161C	25C19	161CA	18.68	161C	25C19	1610	14.83
161C	25C19	162N	15.27	161CA	25C19	1610	29.02
161CA	25C19	162N	28.85	1610	25C19	162N	26.37
660	25C20	67CD1	61.25	660	25C20	163CB	94.46
660	25C20	66C	9.87	660	25C20	68SD	69.49
660	25C20	67CE1	71.63	660	25C20	26CB	36.33
67CD1	25C20	66C	52.72	67CD1	25C20	68SD	78.08
67CD1	25C20	67CE1	17.64	67CD1	25C20	26CB	92.78
163CB	25C20	68 <i>S</i> D	53.55	163CB	25C20	163N	29.80
163CB	25C20	134CB	65.83	163CB	25C20	26CB	58.92
66C	25C20	68SD	74.07	66C	25C20	67CE1	62.03
66C	25C20	26CB	46.18	68SD	25C20	67CE1	95.08
68SD	25C20	163N	82.16	68SD	25C20	134CB	71.67

6000	25020 26-		BLE XVII		
68SD	25C20 26CB	55.82	163N	25C20 134CB	62.79
163N	25C20 26CB	80.06	67CD1	25C21 209CD2	74.52
67CD1	25C21 67CE1	21.15	67CD1	25C21 660	65.47
67CD1	25C21 68SD	94.26	67CD1	25C21 67CA	41.35
67CD1	25C21 67CG	10.50	67CD1	25C21 209CG	78.82
67CD1	25C21 66C	53.86	209CD2	25C21 67CE1	74.15
209CD2	25C21 68SD	93.95	209CD2	25C21 134CB	66.63
209CD2	25C21 67CA	99.15	209CD2	25C21 67CG	80.75
209CD2	25C21 209CG	13.43	209CD2	25C21 68CE	78.53
67CE1	25C21 660	75.62	67CE1	25C21 67CA	60.36
67CE1	25C21 67CG	30.58	67CE1	25C21 209CG	82.83
67CE1	25C21 66C	64.25	660	25C21 68SD	70.26
660	25C21 67CA	36.25	660	25C21 67CG	56.88
660	25C21 68CE	92.80	660	25C21 66C	11.70
68SD	25C21 134CB	81.96	68SD	25C21 67CA	59.58
68SD	25C21 67CG	85.39	68SD	25C21 209CG	80.91
68SD	25C21 68CE	22.58	68SD	25C21 66C	74.79
134CB	25C21 209CG	61.74	134CB	25C21 68CE	60.74
67CA	25C21 67CG	30.85	67CA	25C21 209CG	95.15
67CA	25C21 68CE	78.69	67CA	25C21 66C	29.16
67CG	25C21 209CG	82.94	67CG	25C21 68CE	99.01
67CG	25C21 66C	45.56	209CG	25C21 68CE	65.12
68CE	25C21 66C	97.17	163N	25C22 163CB	41.38
163N	25C22 162C	21.72	163N	25C22 163CA	22.86
163N	25C22 1620	35.18	163N	25C22 162CA	35.98
163N	25C22 162N	54.90	163N	25C22 134CB	83.08
163N	25C22 25SG	48.42	163N	25C22 161C	68.41
163N	25C22 68SD	98.47	163N	25C22 1610	68.27
163N	25C22 134CA	66.14	163CB	25C22 162C	60.85
163CB	25C22 163CA	23.53	163CB	25C22 1620	65.54
163CB	25C22 162CA	77.32	163CB	25C22 162N	96.04
163CB	25C22 134CB	83.53	163CB	25C22 25SG	60.46
1.63CB	25C22 660	95.09	163CB	25C22 68SD	58.15
163CB	25C22 134CA	68.94	162C	25C22 163CA	38.46
162C	25C22 162O	18.52	162C	25C22 162CA	21.19
162C	25C22 162N	35.80	162C	25C22 134CB	73.18
162C	25C22 25SG	61.51	162C	25C22 161C	51.73
162C	25C22 1610	56.33	162C	25C22 134CA	58.55

			TA	BLE XVII			
163CA	25C22	1620	42.12	163CA	25C22	162CA	57.32
163CA	25C22	162N	74.25	163CA	25C22	134CB	72.12
163CA	25C22	25SG	61.97	163CA	25C22	161C	89.59
163CA	25C22	68SD	76.50	163CA	25C22	1610	90.96
163CA	25C22	134CA	55.19	1620	25C22	162CA	34.60
1620	25C22	162N	40.22	1620	25C22	134CB	54.86
1620	25C22	25SG	<b>79.</b> 79	1620	25C22	161C	57.52
1620	25C22	1610	66.63	1620	25C22	134CA	41.20
162CA	25C22	162N	20.16	162CA	25C22	134CB	87.29
162CA	25C22	25 <b>S</b> G	55.65	162CA	25C22	161C	32.48
162CA	25C22	1610	35.17	162CA	25C22	134CA	75.53
162N	25C22	134CB	81.91	162N	25C22	25SG	71.75
162N	25C22	161C	17.31	162N	25C22	1610	28.19
162N	25C22	134CA	75.24	134CB	25C22	161C	94.20
134CB	25C22	68SD	76.86	134CB	25C22	134CA	17.24
25SG	25C22	161C	71.58	25SG	25C22	660	87.43
25SG	25C22	1610	59.61	161C	25C22	1610	14.91
161C	25C22	134CA	90.49	660	25C22	68SD	61.09
68SD	25C22	134CA	77.47	1610	25N23	25\$G	96.69
1610	25N23	162ND1	87.98	1610	25N23	161C	7.60
1610	25N23	162CA	42.67	1610	25N23	162CB	53.10
1610	25N23	162N	24.12	1610	25N23	162CG	72.20
1610	25N23	162CE1	99.02	25SG	25N23	162ND1	58.85
25SG	25N23	161C	90.78	25SG	25 <b>N2</b> 3	162CA	65.10
25 <i>S</i> G	25N23	162CB	74.63	25SG	25N23	25CB	22.39
25SG	25N23	162N	78.71	25SG	25 <b>N2</b> 3	162CG	65.53
25SG	25N23	162CE1	56.82	162ND1	25 <b>N2</b> 3	161C	80.81
162ND1	25N23	162CA	47.69	162ND1	25 <b>N2</b> 3	162CB	35.43
162ND1	25N23	25CB	47.77	162ND1	25 <b>N2</b> 3	162N	64.92
162ND1	25N23	162CG	16.11	162ND1	. 25 <b>N23</b>	162CE1	11.08
161C	25N23	162CA	35.07	161C	25N23	162CB	46.35
161C	25N23	162N	16.53	161C	25 <b>N2</b> 3	162CG	65.21
161C	25N23	162CE1	91.80	162CA	25N23	162CB	20.44
162CA	25N23	25CB	73.45	162CA	25N23	162N	18.55
162CA	25N23	162CG	33.95	162CA	25 <b>N2</b> 3	162CE1	58.16
162CB	25N23	25CB	75.17	162CB	25N23	162N	31.96
162CB	25N23	162CG	19.34	162CB	25 <b>N23</b>	162CE1	46.49
25CB	25 <b>N23</b>	162N	90.62	25CB	25 <b>N2</b> 3	162CG	59.94

		TA	BLE XVII		
25CB	25N23 162CE1	41.59	162N	25N23 162CG	49.81
162N	25N23 162CE1	75.77	162CG	25N23 162CE1	27.14
1610	25C24 162ND1	71.20	1610	25C24 25SG	68.87
1610	25C24 25CB	86.70	1610	25C24 162CE1	84.02
1610	25C24 161C	5.61	1610	25C24 162CG	60.27
2300H2	25C24 19NE2	82.86	2300H2	25C24 23CA	39.21
162ND1	25C24 25SG	49.38	162ND1	25C24 19NE2	75.30
162ND1	25C24 25CB	42.32	162ND1	25C24 162CE1	12.89
162ND1	25C24 161C	66.20	162ND1	25C24 162CG	13.50
25 <b>S</b> G	25C24 19NE2	69.24	25SG	25C24 25CB	21.54
25 <b>S</b> G	25C24 162CE1	51.86	25SG	25C24 161C	67.93
25 <b>S</b> G	25C24 23CA	75.54	25SG	25C24 162CG	55.53
19NE2	25C24 25CB	50.50	19NE2	25C24 162CE1	63.30
19NE2	25C24 23CA	49.37	19NE2	25C24 162CG	88.70
25CB	25C24 162CE1	38.82	25CB	25C24 161C	84.51
25CB	25C24 23CA	72.33	25CB	25C24 162CG	53.33
162CE1	25C24 161C	79.07	162CE1	25C24 162CG	25.60
161C	25C24 162CG	54.95	2300H2	25025 23CA	52.31
2300H2	25025 220	74.89	2300H2	25025 23N	49.81
2300H2	25025 23C	64.37	23CA	25025 19NE2	59.20
23CA	25025 25 <i>S</i> G	80.98	23CA	25025 220	33.94
23CA	25025 23N	14.51	23CA	25025 23C	15.82
19NE2	25025 25SG	68.80	19NE2	25025 220	32.41
19NE2	25025 23N	57.42	19NE2	25025 23C	55.28
25SG	25025 220	87.84	25 <b>SG</b>	25025 1610	56.79
25SG	25025 23N	93.48	25 <b>S</b> G	25025 23C	65.16
220	25025 23N	26.63	2.20	25025 23C	39.45
23N	25025 23C	29.23	162ND1	25C26 1610	70.33
162ND1	25C26 162CG	16.51	162ND1	25C26 184CZ2	61.61
162ND1	25C26 162CB	33.57	162ND1	25C26 162CE1	14.30
162ND1	25C26 184NE1	54.26	1610	25C26 162CG	62.73
1610	25C26 162CB	46.80	1610	25C26 162CE1	84.49
162CG	25C26 184CZ2	58.73	162CG	25C26 162CB	18.91
162CG	25C26 162CE1	27.71	162CG	25C26 184NE1	51.32
184CZ2	25C26 162CB	<b>69</b> .73	184CZ2	25C26 162CE1	53.77
184CZ2	25C26 184NE1	30.35	162C3	25C26 162CE1	46.35
162CB	25C26 184NE1	78.86	162CE1	25C26 184NE1	40.84
1610	25C27 161OD1	53.78	1610	25C27 161C	11.71

TABLE XVII									
1610	25C27	162CB	46.13	1610	25C27	162ND1	62.87		
1610D1	25C27	161C	42.77	1610D1	25C27	162CB	55.70		
1610D1	25C27	162ND1	86.16	161C	25C27	162CB	40.57		
161C	25C27	162ND1	63.10	162CB	25C27	162ND1	30.49		
1610D1	25C28	1370	69.81	1610D1	25C28	1610	54.98		
1610D1	25C28	137C	53.96	1610D1	25C28	137CB	55.92		
1610D1	25C28	162CB	61.51	1610D1	25C28	161CG	11.62		
1610D1	25C28	138N	47.36	1610D1	25C28	161C	42.86		
1370	25C28	137C	15.86	1370	25C28	184CZ2	70.21		
1370	25C28	137CB	37.24	1370	25C28	162CB	85.15		
1370	25C28	161CG	80.00	1370	25C28	138N	27.64		
1370	25C28	184CH2	55.56	1610	25C28	137CB	84.76		
1610	25C28	162CB	45.37	1610	25C28	161CG	49.97		
1610	25C28	161C	13.42	137C	25C28	184CZ2	81.49		
137C	25C28	137CB	32.38	137C	25C28	162CB	78.21		
137C	25C28	161CG	64.18	137C	25C28	138N	15.77		
137C	25C28	184CH2	68.22	137C	25C28	161C	90.09		
184CZ2	25C28	137CB	63.63	184CZ2	25C28	162CB	70.10		
184CZ2	25C28	138N	96.79	184CZ2	25C28	184CH2	15.72		
137CB	25C28	162CB	47.93	137CB	25C28	161CG	67.40		
137CB	25C28	138N	45.63	137CB	25C28	184CH2	56.70		
137CB	25C28	161C	73.01	162CB	25C28	161CG	66.72		
162CB	25C28	138N	87.00	162CB	25C28	184CH2	76.56		
162CB	25C28	161C	39.92	161CG	25C28	138N	55.54		
161CG	25C28	161C	39.76	138N	25C28	184CH2	82.84		
138N	25C28	161C	88.57	1610D1	25C29	161CG	17.77		
1610D1	25C29	161CB	38.20	1610D1	25C29	137C	69.75		
1610D1	25C29	138N	63.36	1610D1	25C29	1610	67.46		
1610D1	25C29	1370	86.87	1610 <b>D1</b>	25C29	138CA	77.48		
1610D1	25C29	137CB	66.56	1610D1	25C29		51.63		
1610D1	25C29	137CA	57.43	1610D1	25C29		38.54		
1610D1	25C29	161ND2	20.78	1610 <b>D1</b>	25C29	162CB	69.24		
1610D1	25C29	138CB	70.48	1610 <b>D1</b>	25C29	161CA	39.44		
1610D1	25C29	162N	49.70	161CG		161CB	23.57		
161CG	25C29	137C	84.96	161CG		138N	74.84		
161CG	25C29	1610	62.41	161CG		138CA	84.52		
161CG	25C29	137CB	84.24	161CG		161C	49.09		
161CG	25 <b>C29</b>	137CA	74.70	161CG	25C29	137N	55.87		

			TA	BLE XVII			•
161CG	25C29	161ND2	11.82	161CG	25C29	162CB	78.89
161CG	25C29	138CB	72.41	161CG	25C29	161CA	31.89
161CG	25C29	162N	53.19	161CB	25C29	138N	98.38
161CB	25C29	1610	43.85	161CB	25C29	137CB	99.44
161CB	25C29	161C	35.43	161CB	25C29	137CA	94.94
161CB	25C29	137N	76.23	161CB	25C29	161ND2	33.15
161CB	25C29	162CB	75.97	161CB	25C29	138CB	91.71
161CB	25C29	161CA	16.99	161CB	25C29	162N	46.28
137C	25C29	138N	19.59	137C	25C29	1370	17.39
137C	25C29	138CA	35.03	137C	25C29	137CB	35.61
137C	25C29	137CA	20.11	137C	25C29	137N	34.13
137C	25C29	161ND2	78.64	137C	25C29	162CB	84.64
137C	25C29	138CB	50.91	137C	25C29	162N	96.62
138N	25C29	1370	32.46	138N	25C29	138CA	20.27
138N	25C29	137CB	52.67	138N	25C29	137CA	33.41
138N	25C29	137N	37.77	138N	25C29	161ND2	65.96
138N	25C29	162CB	98.96	138N	25C29	138CB	32.14
1610	25C29	137CB	93.41	1610	25C29	161C	16.38
1610	25C29	137N	91.81	1610	25C29	161ND2	74.07
1610	25C29	162CB	47.11	1610	25C29	161CA	30.52
1610	25C29	162N	27.28	1370	25C29	138CA	39.55
1370	25C29	137CB	39.61	1370	25C29	137CA	32.50
1370	25C29	137N	49.84	1370	25C29	161ND2	95.83
1370	25C29	162CB	87.74	1370	25C29	138CB	58.44
138CA	25C29	137CB	70.56	138CA	25C29	137CA	52.71
138CA	25C29	137N	57.82	138CA	25C29	161ND2	73.59
138CA	25C29	138CB	19.26	137CB	25C29	161C	82.69
137CB	25C29	137CA	20.23	137CB	25C29	137N	32.22
137CB	25C29	161ND2	84.94	137CB	25C29	162CB	49.44
137CB	25C29	138CB	84.72	137CB	25C29	161CA	89.09
137CB	25C29	162N	67.71	161C	25C29	137CA	89.83
161C	25C29	137N	76.15	151C	25C29	161ND2	60.90
161C	25C29	162CB	43.16	1.61.C	25C29	161CA	18.88
161C	25C29	162N	15.26	137CA	25C29	1.37N	18.89
137CA	25C29	161ND2	71.31	137CA	25C29	162CB	66.00
137CA	25C29	138CB	65.00	137CA	25C29	161CA	89.85
137CA	25C29	162N	76.62	137N	25C29	161ND2	53.90
137N	25C29	162CB	64.20	137N	25C29	138CB	63.74

TABLE XVII									
137N	25C29	161CA	72.67	137N	25C29	162N	65.28		
161ND2	25C29	162CB	88.34	161ND2	25C29	138CB	60.67		
161ND2	25C29	161CA	43.56	161ND2	25C29	162N	64.33		
162CB	25C29	161CA	59.23	162CB	25C29	162N	29.75		
161CA	25C29	162N	29.47	1370	25C30	143NE2	59.46		
1370	25C30	184CZ2	69.76	1370	25C30	137C	14.57		
1370	25C30	138CA	35.79	1370	25C30	184CH2	56.99		
1370	25C30	138N	26.64	143NE2	25C30	184CZ2	83.66		
143NE2	25C30	137C	71.37	143NE2	25C30	138CA	63.07		
143NE2	25C30	184CH2	68.61	143NE2	25C30	138N	72.68		
184CZ2	25C30	137C	78.03	184CZ2	25C30	184CH2	16.45		
184CZ2	25C30	138N	93.46	137C	25C30	138CA	30.17		
137C	25C30	184CH2	67.67	137C	25C30	138N	15.43		
138CA	25C30	184CH2	92.20	138CA	25C30	138N	17.17		
184CH2	25C30	138N	82.69	184CZ2	25N31	184NE1	33.41		
184CZ2	25N31	184CE2	17.32	184CZ2	25N31	162ND1	57.89		
184CZ2	25N31	19NE2	96.35	184NE1	25 <b>N3</b> 1	184CE2	16.98		
184NE1	25N31	162ND1	53.50	184NE1	25N31	19NE2	63.65		
184CE2	25N31	162ND1	57.86	184CE2	25N31	19NE2	80.63		
162ND1	25N31	19NE2	68.39	184NE1	25C32	184CE2	19.92		
184NE1	25C32	184CZ2	38.87	184NE1	25C32	19NE2	81.55		
184NE1	25C32	190E1	51.17	184NE1	25C32	19CD	64.02		
184NE1	25C32	184CD1	13.69	184NE1	25C32	162ND1	58.56		
184NE1	25C32	162CE1	44.43	184CE2	25C32	184CZ2	20.46		
184CE2	25C32	190E1	70.66	184CE2	25C32	19CD	83.94		
184CE2	25C32	184CD1	29.80	184CE2	25C32	162ND1	62.83		
184CE2	25C32	162CE1	52.80	184CZ2	25C32	190E1	86.31		
184CZ2	25C32	184CD1	50.14	184CZ2	25C32	162ND1	60.09		
184CZ2	25C32	162CE1	56.05	19NE2	25C32	190E1	31.34		
19NE2	25C32	19CD	17.91	19NE2	25C32	184CD1	75.00		
19NE2	25C32	162ND1	75.92	19NE2	25C32	162CE1	69.31		
190E1	25C32	19CD	16.54	190E1	25C32	184CD1	47.47		
190E1	25C32	162ND1	56.39	190E1	25C32	162CE1	44.21		
19CD	25C32	184CD1	57.09	19CD	25C32	162ND1	70.79		
19CD	25C32	162CE1	60.09	184CD1	25C32	162ND1	69.23		
184CD1	25C32	162CE1	53.91	162ND1	25C32	162CE1	16.00		
184NE1	25033	190E1	71.37	184NE1	25033		86.43		
184NE1	25033	184CE2	20.79	184NE1	25033	184CD1	17.17		

			TA	BLE XVII			
184NE1	25033	162CE1	60.07	184NE1	25033	184CZ2	41.79
184NE1	25033	162ND1	76.01	184NE1	25033	19CG	80.45
184NE1	25033	184CD2	16.63	184NE1	25033	162NE2	50.59
184NE1	25033	184CG	13.72	184NE1	25033	162CG	72.21
184NE1	25033	25CB	93.48	190E1	25033	19NE2	40.72
190E1	25033	19CD	21.44	190E1	25033	184CE2	91.61
190E1	25033	184CD1	61.45	190E1	25033	162CE1	59.41
190E1	25033	162ND1	74.45	190E1	25033	19CG	30.46
190E1	25033	184CD2	87.26	190E1	25033	162NE2	64.69
190E1	25033	184CG	72.71	190E1	25033	162CG	83.38
190E1	25033	25CB	44.37	19NE2	25033	19CD	22.85
19NE2	25033	184CD1	95.28	19NE2	25033	162CE1	91.27
19NE2	25033	162ND1	97.51	19NE2	25033	19CG	32.33
19NE2	25033	25CB	55.21	19CD	25033	184CD1	72.73
19CD	25033	162CE1	79.91	19CD	25033	162ND1	92.65
19CD	25033	19CG	15.53	19CD	25033	162NE2	85.98
19CD	25033	184CG	83.64	19CD	25033	25CB	55.10
184CE2	25033	184CD1	35.82	184CE2	25033	162CE1	66.09
184CE2	25033	184CZ2	21.88	184CE2	25033	162ND1	76.32
184CE2	25033	184CD2	10.76	184CE2	25033	162NE2	54.99
184CE2	25033	184CG	26.75	184CE2	25033	162CG	68.94
184CD1	25033	162CE1	68.79	184CD1	25033	184CZ2	57.65
184CD1	25033	162ND1	87.29	184CD1	25033	19CG	64.62
184CD1	25033	184CD2	27.61	184CD1	25033	162NE2	61.38
184CD1	25033	184CG	11.29	184CD1	25033	162CG	85.74
184CD1	25033	25CB	93.60	162CE1	25033	184CZ2	67.50
162CE1	25033	162ND1	20.07	162CE1	25033	19CG	89.40
162CE1	25033	184CD2	72.00	162CE1	25033	162NE2	11.22
162CE1	25033	184CG	72.92	162CE1	25033	162CG	25.06
162CE1	25033	25CB	41.49	184CZ2	25033	162ND1	70.13
184CZ2	25033	184CD2	31.61	184CZ2	25033	162NE2	56.74
184CZ2	25033	184CG	48.49	184CZ2	25033	162CG	60.18
162ND1	25033	184CD2	84.61	162ND1	25033	162NE2	25.95
162ND1	25033	184CG	89.59	1.62ND1	25033	162CG	11.10
162ND1	25033	25CB	42.38	19CG	25033	184CD2	91.76
19CG	25033	162NE2	93.17	19 <b>CG</b>	25033	184CG	74.56
19CG	25033	25CB	70.08	184CD2	25033	162NE2	61.34
184CD2	25033	184CG	17.21	184CD2	25033	162CG	78.20

			TA	BLE XVII			
162NE2	25033 1	L84CG	63.94	162NE2	25033	162CG	26.05
162NE2	25033	25CB	52.56	184CG	25033	162CG	85.93
162CG	25033	25CB	53.49	184NE1	25034	184CE2	18.11
184NE1	25034 1	L84CZ2	33.70	184NE1	25034	19NE2	70.03
184NE1	25034 1	184CD1	15.57	184NE1	25034	19CD	56.05
184NE1	25034	190E1	43.11	184CE2	25034	184CZ2	18.11
184CE2	25034	19NE2	87.63	184CE2	25034	184CD1	28.12
184CE2	25034	19CD	74.10	184CE2	25034	190E1	60.83
184CZ2	25034	19NE2	97.82	184CZ2	25034	184CD1	46.02
184CZ2	25034	19CD	86.79	184CZ2	25034	190E1	72.54
19NE2	25034 1	L84CD1	68.22	19NE2	25034	19CD	16.05
19NE2	25034	190E1	26.92	184CD1	25034	19CD	52.52
184CD1	25034	190E1	42.67	19CD	25034	190E1	14.42
200	25C35	19NE2	68.42	200	25C35	19CD	70.46
200	25C35	19CG	55.17	200	25C35	190E1	84.98
200	25C35	21NE2	59.84	200	25C35	210E1	59.50
19NE2	25C35 1	184NE1	74.82	19NE2	25C35	19CD	18.63
19NE2	25C35	19CG	33.10	19NE2	25C35	184CD1	76.03
19NE2	25C35	190E1	29.73	19NE2	25C35	184CE2	89.27
184NE1	25C35	19CD	61.33	184NE1	25C35	19CG	69.26
184NE1	25C35 1	L84CD1	18.20	184NE1	25C35	190E1	46.19
184NE1	25C35 1	L84CE2	15.64	19CD	25C35	19CG	20.18
19CD	25C35 1	L84CD1	59.00	19CD	25C35	190E1	15.55
19CD	25C35 1	L84CE2	76.79	19CG	25C35	184CD1	60.31
19CG	25C35	190E1	31.00	19CG	25C35	184CE2	84.62
184CD1	25C35	19021	46.45	184CD1	25 <b>C</b> 35	184CE2	27.90
190E1	25C35 1	L84CE2	61.48	21NE2	25C35	210E1	26.73
21NE2	25C36	200	62.73	21NE2	25C36	21CD	14.08
200	25C36	19CG	49.13	200	25C36	21CD	53.84
184CD1	25C36 1	184NE1	18.40	184CD1	25C36	184CG	16.42
184CD1	25C36 1	L84CE2	27.92	184CD1	25C36	19CG	56.22
184NE1	25C36 1	L84CG	28.30	184NE1	25C36	184CE2	16.44
184NE1	25C36	19CG	61.95	184CG	25C36	184CE2	27.91
184CG	25C36	19CG	70.47	184CE2	25C36	19CG	78.31
21NE2	25C37	200	79.74	21NE2	25C37	21CD	16.72
21NE2	25C37	210E1	29.73	21NE2	25C37	20C	72.88
200	25C37	21CD	66.28	200	25C37	210E1	66.79
200	25C37	20C	14.00	21CD	25C37	210E1	16.17

			TA	BLE XVII			
21CD	25C37	20C	62.16	210E1	25C37	20C	66.88
21NE2	25C38	21CD	9.90	21NE2	25C38	200	63.92
21CD	25C38	200	54.43	184CG	25C40	184CB	20.90
184CG	25C40	184CD2	19.97	184CG	25C40	184CD1	18.76
184CG	25C40	184CE2	29.83	184CG	25C40	184NE1	29.07
184CG	25C40	184CE3	34.09	184CG	25C40	1840	55.54
184CG	25C40	184CA	32.20	184CB	25C40	184CD2	36.24
184CB	25C40	184CD1	35.47	184CB	25C40	184CE2	50.10
184CB	25C40	184NE1	49.38	184CB	25C40	184CE3	43.19
184CB	25C40	1840	34.81	184CB	25C40	184CA	17.73
184CD2	25C40	184CD1	30.80	184CD2	25C40	184CE2	17.77
184CD2	25C40	184NE1	29.02	184CD2	25C40	184CE3	17.34
184CD2	25C40	1840	69.94	184CD2	25 <b>C4</b> 0	184CA	51.16
184CD1	25C40	184CE2	28.86	184CD1	25C40	184NE1	17.14
184CD1	25C40	184CE3	47.95	184CD1	25C40	1840	66.59
184CD1	25C40	184CA	38.47	184CE2	25C40	184NE1	17.07
184CE2	25C40	184CE3	30.61	184CE2	25C40	1840	84.89
184CE2	25C40	184CA	61.54	184NE1	25C40	184CE3	45.35
184NE1	25C40	1840	82.69	184NE1	25C40	184CA	55.30
184CE3	25C40	1840	71.49	184CE3	25C40	184CA	60.55
1840	25C40	184CA	28.85	184CD1	25C41	184CG	21.77
184CD1	25C41	184NE1	21.97	184CD1	25C41	184CD2	35.07
184CD1	25C41	184CE2	34.98	184CD1	25C41	184CB	36.36
184CD1	25C41	184CE3	51.22	184CD1	25C41	184CZ2	50.86
184CD1	25C41	184CA	37.30	184CG	25C41	184NE1	35.87
184CG	25C41	184CD2	21.86	184CG	25C41	184CE2	35.60
184CG	25C41	184CB	19.28	184CG	25C41	184CE3	34.71
184CG	25C41	184CZ2	50.97	184CG	25C41	184CA	30.15
184NE1	· 25C41	184CD2	34.95	184NE1	25C41	184CE2	20.95
184NE1	25C41	184CB	54.49	184NE1	25C41	184CE3	50.02
184NE1	25C41	184CZ2	33.28	184NE1	25C41	184CA	58.90
184CD2	25C41	184CE2	21.45	184CD2	25C41	184CB	36.52
184CD2	25C41	184CE3	16.39	184CD2	25C41	184CZ2	32.78
184CD2	25C41	184CA	51.13	184CE2	25C41	184CB	54.20
		184CE3				184CZ2	
184CE2	25C41	184CA	65.12	184CB	25 <b>C41</b>	184CE3	43.07
184CB	25C41	184CZ2	68.63	184CB	25C41	184CA	17.27
184CE3	25C41	184CZ2	36.53	184CE3	25C41	184CA	59.94

			TAB	LE XVII			
184CZ2	25C41	184CA	80.95	25SG	25042	25CB	38.64
25SG	25042	25N	73.56	25SG	25042	25CA	53.59
25SG	25042	24C	84.78	25SG	25042	190E1	88.63
25 <b>S</b> G	25042	25C	44.83	25SG	25042	26N	45.60
25SG	25042	162ND1	49.49	25 <i>S</i> G	25042	162CE1	54.51
25SG	25042	26CD1	74.03	25CB	25042	25N	49.39
25CB	25042	19NE2	81.36	25CB	25042	24N	99.21
25CB	25042	25CA	25.75	25CB	25042	24C	62.76
25CB	25042	24CA	84.14	25CB	25042	19CD	66.86
25CB	25042	190E1	53.70	25CB	25042	25C	31.71
25CB	25042	26N	46.47	25CB	25042	162ND1	47.29
25CB	25042	162CE1	39.19	25CB	25042	26CD1	82.79
25N	25042	23C	70.38	25N	25042	23CA	94.53
25N	25042	19NE2	73.90	25N	25042	24N	51.16
25N	25042	25CA	23.79	25N	25042	230	72.18
25N	25042	24C	13.37	25N	25042	24CA	34.78
25N	25042	19CD	67.53	25N	25042	190E1	67.99
25N	25042	220	80.77	25N	25042	25C	28.88
25N	25042	26N	36.19	25N	25042	162ND1	95.82
25N	25042	23N	91.99	25N	25042	22C	85.72
25N	25042	162CE1	84.41	25N	25042	26CD1	52.11
23C	25042	23CA	27.54	23C	25042	19NE2	81.75
23C	25042	24N	23.31	23C	25042	25CA	94.12
23C	25042	230	20.23	23C	25042	24C	57.01
23C	25042	24CA	35.64	23C	25042	19CD	92.20
23C	25042	220	49.02	23C	25042	2300H2	70.61
23C	25042	25C	93.36	23C	25042	26N	86.51
23C	25042	23N	31.08	23C	25042	22C	39.18
23C	25042	26CD1	57.51	23CA	25042	19NE2	75.36
23 <b>CA</b>	25042	24N	43.40	23CA	25042	230	40.93
23CA	25042	24C	81.51	23CA	25042	24CA	60.46
23CA	25042	19CD	89.50	23CA	25042	220	38.60
23CA	25042	2300Н2	46.00	23CA	25042	23N	9.62
23CA	25042	22C	25.11	23CA	25042	26CD1	82.64
19NE2	25042	24N	65.75	19NE2	25042	25CA	78.16
19NE2	25042	24C	73.53	19NE2	25042	24CA	73.72
19NE2	25042	19CD	14.86	19NE2	25042	190E1	30.78
19NE2	25042	220	36.84	19NE2	25042	2300H2	94.24

			TA	BLE XVII			
19NE2	25042	25C	94.02	19NE2	25042	162ND1	84.58
19NE2	25042	23N	65.76	19NE2	25042	22C	50.27
19NE2	25042	162CE1	71.88	24N	25042	25CA	74.59
24N	25042	230	38.03	24N	25042	24C	38.35
24N	25042	24CA	18.53	24N	25042	19CD	72.99
24N	25042	190E1	86.24	24N	25042	220	42.70
24N	25042	2300H2	89.40	24N	25042	25C	78.15
24N	25042	26N	76.91	24N	25042	23N	41.41
24N	25042	22C	39.81	24N	25042	26CD1	59.70
25CA	25042	230	93.85	25CA	25042	24C	37.14
25CA	25042	24CA	58.56	25CA	25042	19CD	66.70
25CA	25042	190E1	60.00	25CA	25042	220	97.56
25CA	25042	25C	16.18	25CA	25042	26N	32.48
25CA	25042	162ND1	72.81	25CA	25042	162CE1	62.96
25CA	25042	26CD1	63.86	230	25042	24C	59.84
230	25042	24CA	41.50	230	25042	220	69.15
230	25042	2300H2	73.61	230	25042	25C	87.62
230	25042	26N	76.22	230	25042	23N	47.69
230	25042	22C	58.56	230	25042	26CD1	41.93
24C	25042	24CA	21.42	24C	25042	19CD	70.59
24C	25042	190E1	74.79	24C	25042	220	72.25
24C	25042	25C	40.11	24C	25042	26N	42.88
24C	25042	23N	79.63	24C	25042	22C	75.07
24C	25042	162CE1	96.91	24C	25042	26CD1	47.33
24CA	25042	19CD	76.47	24CA	25042	190E1	85.89
24CA	25042	220	59.12	24CA	25042	25C	60.08
24CA	25042	26N	58.46	24CA	25042	23N	59.62
24CA	25042	22C	58.07	24CA	25042	26CD1	46.49
19CD	25042	190E1	16.46	19CD	25042	220	50.92
19CD	25042	25C	82.87	19CD	25042	26N	98.50
19CD	25042	162ND1	72.56	19CD	25042	23N	79.96
19CD	25042	22C	64.41	19CD	25042	162CE1	58.81
190E1	25042	220	67.33	190 <b>E1</b>	25042	25C	75.67
190E1	25042	26N	92.41	190E1	25042	162ND1	56.82
190E1	25042	23N	96.37	190E1	25042	22C	80.82
190E1	25042	162CE1	42.61	220	25042	2300H2	68.87
220	25042	23N	29.05	220	25042	22C	13.49
2300H2	25042	23N	49.64	2300H2	25042	22C	59.44

			TA	BLE XVII			
25C	25042	26N	16.85	25C	25042	162ND1	77.20
25C	25042	162CE1	70.85	25C	25042	26CD1	51.53
26N	25042	162ND1	87.91	26N	25042	162CE1	84.47
26N	25042	26CD1	36.52	162ND1	25042	162CE1	15.10
23N	25042	22C	15.56	23N	25042	26CD1	88.32
22C	25042	26CD1	95.66				

## TABLE XVIII

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
2420H2	25C1	180D1	67.89	2420H2	25C1	18CG	56.00
2420H2	25C1	18ND2	40.01	2420H2	25C1	1840	81.97
2420H2	25C1	184C	88.17	2420H2	25C1	21NE2	62.78
2420H2	25C1	20N	79.06	180D1	25C1	184CD1	79.40
180D1	25C1	184CB	74.81	180D1	25C1	184CG	85.32
180D1	25C1	184CA	53.47	180D1	25C1	18CG	11.94
180D1	25C1	18ND2	28.26	180D1	25C1	184NE1	92.34
180D1	25C1	1840	58.92	180D1	25C1	184C	49.03
180D1	25C1	21NE2	93.14	180D1	25C1	20N	41.90
184CD1	25C1	184CB	38.10	184CD1	25C1	184CG	19.94
184CD1	25C1	184CA	42.92	184CD1	25C1	18CG	90.87
184CD1	25C1	184NE1	15.56	184CD1	25C1	1840	71.50
184CD1	25C1	184CD2	27.97	184CD1	25C1	184C	61.10
184CD1	25C1	20N	84.09	184CB	25C1	184CG	21.98
184CB	25C1	184CA	21.35	184CB	25C1	18CG	81.80
184CB	25C1	18ND2	96.18	184CB	25C1	184NE1	50.19
184CB	25C1	1840	36.35	184CB	25C1	184CD2	33.73
184CB	25C1	184C	32.01	184CG	25C1	184CA	36.19
184CG	25C1	18CG	95.04	184CG	25C1	184NE1	28.76
184CG	25C1	1840	58.26	184CG	25C1	184CD2	15.95
184CG	25C1	184C	51.91	184CA	25C1	18CG	60.97
184CA	25C1	18ND2	76.21	184CA	25C1	184NE1	58.19
184CA	25C1	1840	30.82	184CA	25C1	184CD2	51.40
184CA	25C1	184C	18.27	184CA	25C1	20N	84.75
18CG	25C1	18ND2	16.83	18CG	25C1	1840	59.45
18CG	25C1	184C	52.65	18CG	25C1	21NE2	87.86
18CG	25C1	20N	45.88	18ND2	25C1	1840	68.25
18ND2	25C1	134C	65.08	18ND2	25C1	21NE2	75.52
18ND2	25C1	20N	49.20	184NE1	25C1	1840	85.54
184NE1	25 <b>C</b> 1	184CD2	27.58	134NE1	25C1	184C	76.21
184NE1	25C1	20N	87.93	1840	25C1	184CD2	69.09
1840	25C1	184C	14.96	184CD2	25C1	184C	65.57
184C	25C1	20N	87.94	21NE2	25C1	20N	61.73
184CD1	25C2	184CG	20.80	184CD1	25C2	184NE1	19.39
184CD1	25C2	184CD2	32.31	J.84CD1	25C2	184CB	37.15

			TAE	BLE XVIII			
184CD1	25C2	184CE2	30.89	184CD1	25C2	180D1	66.60
184CD1	25C2	184CA	37.97	184CD1	25C2	184CE3	47.14
184CG	25C2	184NE1	32.65	184CG	25C2	184CD2	20.10
184CG	25C2	184CB	20.97	184CG	25C2	184CE2	31.76
184CG	25C2	180D1	71.61	184CG	25C2	184CA	31.32
184CG	25C2	184CE3	31.69	184NE1	25C2	184CD2	31.41
184NE1	25C2	184CB	52.95	184NE1	25C2	184CE2	18.36
184NE1	25C2	180D1	83.70	184NE1	25C2	184CA	57.03
184NE1	25C2	184CE3	45.36	184CD2	25C2	184CB	36.72
184CD2	25C2	184CE2	18.93	184CD2	25C2	180D1	91.69
184CD2	25C2	184CA	50.71	184CD2	25C2	184CE3	15.04
184CB	25C2	184CE2	52.15	184CB	25C2	2420H2	91.69
184CB	25C2	180D1	60.47	184CB	25C2	184CA	17.64
184CB	25C2	184CE3	41.82	184CE2	25C2	180D1	97.37
184CE2	25C2	184CA	62.35	184CE2	25C2	184CE3	29.53
2420H2	25C2	180D1	48.28	2420H2	25C2	184CA	79.74
180D1	25C2	184CA	43.08	184CA	25C2	184CE3	58.53
184NE1	25C3	184CD1	20.69	184NE1	25C3	184CE2	18.79
184NE1	25C3	184CG	31.66	184NE1	25C3	184CD2	29.94
184NE1	25C3	184CZ2	30.27	184CD1	25C3	184CE2	31.79
184CD1	25C3	184CG	18.43	184CD1	25 <b>C</b> 3	184CD2	29.94
184CD1	25C3	200	97.66	184CD1	25C3	184CZ2	46.78
184CE2	25C3	184CG	31.39	184CE2	25C3	184CD2	18.34
184CE2	25C3	184CZ2	15.37	184CG	25C3	184CD2	18.68
184CG	25C3	184CZ2	46.08	184CD2	25C3	184CZ2	30.03
200	25C3	21NE2	47.07	200	25C4	19CG	63.08
200	25C4	20C	13.93	200	25C4	21NE2	58.24
200	25C4	20N	38.94	200	25C4	19CD	73.49
200	25C4	20CA	29.58	200	25C4	210E1	64.57
200	25C4	180D1	80.18	200	25C4	21CD	55.60
200	25C4	19NE2	66.42	200	25C4	21N	18.83
19CG	25C4	20C	69.89	19CG	25C4	184CD1	59.97
19CG	. 25C4	184NE1	63.38	19CG	25C4	20N	48.04
19CG	25C4	19CD	19.27	19CG	25C4	20CA	64.22
19CG	25C4	184CG	73.51	19CG	25C4	180D1	63.24
19CG	25C4	184CE2	77.80	19CG	25C4	19NE2	28.95
19CG	25C4	21N	80.44	20C	25C4	21NE2	47.51
20C	25 <b>C4</b>	20N	34.03	20C	25C4	19CD	83.36
20C	25C4	20CA	19.08	20C	25C4	210E1	60.05
20C	25C4	180D1	71.76	20C	25C4	21CD	48.39
20C	25C4	19NE2	78.43	20C	25C4	21N	11.83
184CD1	25C4	184NE1	19.36	184CD1	25C4	20N	90.20
184CD1	25C4	19CD	56.87	184CD1	25C4	184CG	14.62
184CD1	25C4	180D1	63.59	184CD1	25C4	184CE2	26.99

			TAI	BLE XVIII			
184CD1	25C4	19NE2	70.60	184NE1	25C4	19CD	53.18
184NE1	25C4	184CG	27.61	184NE1	25C4	180D1	82.57
184NE1	25C4	184CE2	14.41	184NE1	25C4	19NE2	63.56
21NE2	25C4	20N	70.04	21NE2	25C4	20CA	52.20
21NE2	25C4	210E1	28.00	21NE2	25C4	180D1	83.21
21NE2	25C4	21CD	15.16	21NE2	25C4	21N	39.44
20N	25C4	19CD	66.67	20N	25C4	20CA	18.60
20N	25C4	210E1	91.11	20N	25C4	184CG	96.83
20N	25C4	180D1	41.94	20N	25C4	21CD	77.35
20N	25C4	19NE2	70.35	20N	25C4	21N	45.53
19CD	25C4	20CA	81.50	19CD	25C4	184CG	71.48
19CD	25C4	180D1	79.99	19CD	25C4	184CE2	66.85
19CD	25C4	19NE2	15.24	19CD	25C4	21N	92.20
20CA	25C4	210E1	72.56	20CA	25C4	180D1	52.69
20CA	25C4	21CD	58.76	20CA	25C4	19NE2	81.60
20CA	25C4	21N	28.99	210E1	25C4	21CD	14.61
210E1	25C4	21N	48.49	184CG	25C4	180D1	62.90
184CG	25C4	184CE2	27.32	184CG	25C4	19NE2	85.13
180D1	25C4	184CE2	89.23	180D1	25C4	21CD	97.26
180D1	25C4	19NE2	92.18	180D1	25C4	21N	80.98
184CE2	25C4	19NE2	75.84	21CD	25C4	21N	37.64
19NE2	25C4	21N	85.10	200	25C5	20N	55.48
200	25C5	20C	21.07	200	25C5	20CA	44.11
200	25C5	19CG	71.84	200	25C5	21NE2	66.85
200	25C5	19C	51.82	200	25C5	21N	24.93
200	25C5	19CD	75.35	200	25C5	19N	84.18
200	25C5	19CA	67.21	200	25C5	19CB	67.59
200	25C5	21CD	58.45	200	25C5	190	39.96
200	25C5	210E1	62.69	2 <b>0N</b>	25C5	20C	45.30
20N	25C5	20CA	25.13	20N	25C5	180D1	58.18
20N	25C5	19CG	59.77	20N	25C5	21NE2	88.80
20N	25C5	19C	15.04	20N	25C5	21N	55.27
20N	25C5	2420H2	88.85	20N	25C5	18CG	53.24
20N	25C5	19CD	75.42	20N	25C5	19N	36.29
20N	25C5	19CA	29.03	20N	25C5	19CB	45.65
20N	25C5	21CD	91.99	20N	25C5	18ND2	55.37
20N	25C5	190	20.80	20N	25C5	1830	72.28
20C	25C5	20CA	26.08	20C	25C5	180D1	98.89
20C	25C5	19CG	82.87	20C	25C5	21NE2	55.09
20C	25C5	19C	48.39	200	25C5	21N	11.08
20C	25C5	2420H2	97.88	20C	25C5	18CG	90.40
20C	25C5	19CD	90.83	20C	25C5	19N	80.24
20C	25C5	19CA	66.78	20C	25C5	19CB	74.20
20C	25C5	21CD	51.59	26C	25 <b>C</b> 5	18ND2	83.45

			TAE	LE XVIII			
20C	25C5	190	38.43	20C	25C5	210E1	61.08
20CA	25C5	180D1	72.87	20CA	25C5	19CG	80.41
20CA	25C5	21NE2	63.95	20CA	25C5	19C	35.76
20CA	25C5	21N	33.35	20CA	25C5	242OH2	81.51
20CA	25C5	18CG	64.36	20CA	25C5	19CD	93.94
20CA	25C5	19N	61.13	20CA	25C5	19CA	53.26
20CA	25C5	19CB	67.53	20CA	25C5	21CD	66.97
20CA	25C5	18ND2	58.72	20CA	25C5	190	32.23
20CA	25C5	1830	97.26	20CA	25C5	210E1	79.78
180D1	25C5	19CG	80.27	180D1	25C5	19C	65.37
180D1	25C5	184CD1	74.17	180D1	25C5	2420H2	54.74
180D1	25C5	18CG	10.85	180D1	25C5	19CD	93.30
180D1	25C5	184NE1	90.85	180D1	25C5	19N	40.79
180D1	25C5	19CA	57.69	180D1	25C5	19CB	70.95
180D1	25C5	18ND2	26.15	180D1	25 <b>C</b> 5	190	76.52
180D1	25C5	184CG	70.18	180D1	25C5	1830	49.64
19CG	25C5	19C	45.54	19CG	25C5	184CD1	61.26
19CG	25C5	21N	92.85	19CG	25C5	18CG	85.81
19CG	25C5	19CD	16.55	19CG	25C5	184NE1	60.42
19CG	25C5	19N	42.09	19CG	25C5	19CA	32.92
19CG	25C5	19CB	14.39	19CG	25C5	18ND2	99.14
19CG	25C5	190	48.69	19CG	25C5	184CG	74.77
19CG	25C5	1830	42.29	21NE2	25C5	19C	99.03
21NE2	25C5	21N	44.87	21NE2	25C5	2420H2	63.97
21NE2	25C5	18CG	97.62	21NE2	25C5	21CD	14.03
21NE2	25C5	18ND2	81.87	21NE2	25C5	190	91.86
21NE2	25C5	210E1	25.97	19C	25C5	184CD1	99.04
19C	25C5	21N	59.38	19C	25C5	18CG	62.98
19C	25C5	19CD	60.71	19C	25C5	19N	32.80
19C	25C5	19CA	18.65	19C	25C5	19CB	31.95
19C	25C5	21CD	99.25	19C	25C5	18ND2	68.17
19C	25C5	190	11.87	19C	25C5	1830	65.75
184CD1	25C5	18CG	85.01	184CD1	25C5	19CD	55.30
184CD1	25C5	184NE1	16.94	184CD1	25C5	19N	71.72
184CD1	25C5	19CA	80.51	184CD1	25C5	19CB	70.19
184CD1	25C5	18ND2	99.06	184CD1	25C5	184CG	14.35
184CD1	25C5	1830	35.61	21N	25C5	2420H2	94.26
21N	25C5	18CG	95.93	2 J.N	25C5	19CD	99.24
21N	25C5	19N	90.87	21N	25C5	19CA	77.83
21N	25C5	19CB	84.95	21N	25C5	21CD	40.54
21.N	25C5	18ND2	86.56	21N	25C5	190	49.51
21N	25C5	210E1	50.14	2420H2	25C5	18CG	48.98
2420H2	25C5	19N	93.10	2420H2	25C5	21CD	77.70
2420H2	25C5	18ND2	37.02	2420H2	25C5	184CG	92.59

TABLE XVIII									
2420H2	25C5	210E1	85.45	18CG	25C5	19N	44.21		
18CG	25C5	19CA	59.10	18CG	25C5	19CB	74.72		
18CG	25C5	18ND2	15.95	18CG	25C5	190	73.12		
18CG	25C5	184CG	80.57	18CG	25C5	1830	59.45		
19CD	25C5	184NE1	49.55	19CD	25C5	19N	57.74		
19CD	25C5	19CA	49.44	19CD	25C5	19CB	30.83		
19CD	25C5	190	61.72	19CD	25C5	184CG	69.64		
19CD	25C5	1830	48.36	184NE1	25C5	19N	82.36		
184NE1	25C5	19CA	86.84	184NE1	25C5	19CB	72.47		
184NE1	25C5	184CG	26.82	184NE1	25C5	1830	47.92		
19N	25C5	19CA	18.06	19N	25C5	19CB	30.56		
19N	25C5	18ND2	57.06	19N	25C5	190	44.42		
19N	25C5	184CG	78.07	19N	25C5	1830	36.14		
19CA	25C5	19CB	18.64	19CA	25C5	18ND2	69.26		
19CA	25C5	190	28.69	19CA	25C5	184CG	89.98		
19CA	25C5	1830	47.24	19CB	25C5	18ND2	86.66		
19CB	25C5	190	37.11	19CB	25C5	184CG	82.34		
19CB	25C5	1830	43.36	21CD	25C5	18ND2	94.92		
21CD	25C5	190	90.00	21CD	25C5	210E1	14.31		
18ND2	25C5	190	76.17	18ND2	25C5	184CG	92.52		
18ND2	25C5	1830	75.40	190	25C5	1830	75.72		
190	25C5	210E1	98.98	184CG	25C5	1830	42.87		
180D1	25C6	2420H2	77.61	180D1	25C6	20N	62.12		
180D1	25C6	18CG	14.65	180D1	25C6	18ND2	34.76		
180D1	25C6	20CA	76.78	180D1	25C6	184CD1	84.66		
180D1	25C6	20C	97.24	180D1	25C6	19CG	77.43		
180D1	25C6	184CA	51.93	180D1	25C6	184CG	83.75		
180D1	25C6	184CB	70.76	180D1	25C6	19N	38.61		
180D1	25C6	19C	62.85	180D1	25C6	1830	51.93		
180D1	25C6	2440H2	87.25	180D1	25C6	184NE1	97.99		
2420H2	25 <b>C</b> 6	18CG	65.17	2420H2	25C6	18ND2	48.30		
2420H2	25C6	20CA	92.93	2420H2	25C6	21NE2	71.65		
2420H2	25C6	184CA	<b>99.9</b> 9	2420H2	25C6	2440H2	54.20		
20N	25C6	18CG	61.40	20N	25C6	18ND2	64.85		
20N	25C6	20CA	22.20	20N	25C6	200	43.67		
20N	25C6	20C	36.81	20N	25C6	21NE2	80.01		
20N	25C6	19CG	49.35	20N	25C6	19N	37.08		
20N	25C6	19C	10.50	2 <b>0N</b>	25C6	1830	71.86		
20N	25C6	2440H2	62.38	18CG	25 <b>C</b> 6	18ND2	20.14		
18CG	25C6	20CA	71.10	18CG	25C6	184CD1	99.10		
18CG	25C6	20C	92.50	18CG	25C6	19CG	87.70		
18CG	25C6	184CA	64.20	18CG	25C6	184CG	96.81		
18CG	25C8	184CB	82.01	18CG	25C6	19N	47.68		
18CG	25C6	19C	65.05	18CG	25C6	1830	66.43		

			TAI	BLE XVIII			
18CG	25C6	2440H2	73.82	18ND2	25C6	20CA	66.38
18ND2	25C6	20C	86.96	18ND2	25C6	21NE2	91.35
18ND2	25C6	184CA	81.21	18ND2	25C6	184CB	96.97
18ND2	25C6	19N	63.62	18ND2	25C6	19C	71.84
18ND2	25C6	1830	86.52	18ND2	25C6	2440H2	56.34
20CA	25C6	200	36.02	20CA	25C6	20C	21.45
20CA	25C6	21NE2	58.02	20CA	25C6	19CG	66.36
20CA	25C6	19N	58.93	20CA	25C6	19C	31.05
20CA	25C6	1830	94.02	20CA	25C6	2440H2	42.94
200	25C6	20C	17.19	200	25C6	21NE2	55.96
200	25C6	19CG	53.96	200	25C6	19N	73.45
200	25C6	19C	43.47	200	25C6	1830	95.13
200	25C6	2440H2	65.82	200	25C6	184NE1	94.22
184CD1	25C6	19CG	55.80	184CD1	25C6	184CA	40.42
184CD1	25C6	184CG	17.20	184CD1	25C6	184CB	33.57
184CD1	25C6	19N	72.31	184CD1	25C6	19C	91.74
184CD1	25C6	1830	36.78	184CD1	25C6	184NE1	14.58
20C	25C6	21NE2	47.89	20C	25C6	19CG	64.56
20C	25C6	19N	72.48	20C	25C6	19C	41.10
20C	25C6	2440H2	49.53	21NE2	25C6	19C	87.32
21NE2	25C6	2440H2	35.45	19CG	25C6	184CA	76.21
19CG	25C6	184CG	72.47	19CG	25C6	184CB	83.57
19CG	25C6	19N	40.03	19CG	25C6	19C	38.92
19CG	25C6	1830	42.08	19CG	25C6	184NE1	53.94
184CA	25C6	184CG	32.82	184CA	25C6	184CB	19.38
184CA	25C6	19N	63.55	184CA	25C6	19C	94.44
184CA	25C6	1830	34.70	184CA	25C6	184NE1	54.88
184CG	25C6	184CB	18.90	184CG	25C6	19N	82.27
184CG	25C6	1830	45.36	184CG	25C6	184NE1	27.16
184CB	25C6	19N	80.80	184CB	25C6	1830	46.93
184CB	25C6	184NE1	45.74	19N	25C6	19C	31.43
19N	25C6	1830	36.92	19N	25C6	2440H2	93.07
19N	25C6	184NE1	79.67	19C	25C6	1830	63.42
19C	25C6	2440H2	72.61	19C	25C6	184NE1	92.57
1830	25C6	184NE1	47.52	200	25C7	19CG	66.72
200	25C7	20C	8.16	200	25C7	19CD	83.83
200	25C7	19ME2	82.44	200	25C7	220	56.54
200	25C7	22N	35.30	20C	25C7	21CA	31.83
200	25C7	190E1	95.92	200	25C7	210E1	66.65
200	25C7	21N	17.17	200	25C7	20N	30.97
200	25C7	21NE2	51.64	200	25C7	20CA	19.21
19CG	25C7	20C	71.45	19CG	25C7	19CD	21.67
19CG	25C7	19NE2	34.92	19CG	25 <b>C7</b>	220	55.96
19CG	25C7	184NE1	60.44	19CG	25C7	22N	79.46

			TAI	BLE XVIII			
19CG	25C7	21CA	96.64	19CG	25C7	190E1	30.14
19CG	25C7	21N	83.86	19CG	25C7	184CD1	53.93
19CG	25C7	20N	45.16	19CG	25C7	20CA	61.57
20C	25C7	19CD	89.91	20C	25C7	19NE2	89.95
20C	25C7	220	64.63	20C	25C7	22N	42.08
20C	25C7	21CA	30.95	20C	25C7	210E1	60.76
20C	25C7	21N	13.74	20C	25C7	20N	31.22
20C	25C7	21NE2	43.84	20C	25C7	20CA	15.80
19CD	25C7	19NE2	18.60	19CD	25C7	220	53.43
19CD	25C7	184NE1	53.45	19CD	25C7	22N	85.40
19CD	25C7	190E1	13.55	19CD	25C7	184CD1	54.25
19CD	25C7	20N	66.44	19CD	25C7	20CA	82.18
19NE2	25C7	220	38.61	19NE2	25C7	184NE1	67.89
19NE2	25C7	22N	73.27	19NE2	25C7	190E1	28.20
19NE2	25C7	21N	97.49	19NE2	25C7	184CD1	71.68
19NE2	25C7	20N	74.18	19NE2	25 <b>C7</b>	20CA	86.96
220	25C7	22N	35.17	220	25C7	21CA	64.61
220	25 <b>C</b> 7	190E1	65.85	220	25C7	21N	66.20
220	25C7	20N	67.78	220	25C7	20CA	70.65
184NE1	25C7	190E1	40.45	184NE1	25C7	184CD1	16.51
184NE1	25C7	20N	91.81	22N	25C7	21CA	29.87
22N	25C7	190E1	98.77	22N	25C7	210E1	70.75
22N	25C7	21N	36.58	22N	25C7	20N	63.54
22N	25C7	21NE2	72.49	22N	25C7	20CA	55.44
21CA	25C7	210E1	42.69	21CA	25C7	21N	17.64
21CA	25C7	20N	61.95	21CA	25C7	21NE2	43.40
21CA	25C7	20CA	46.53	190E1	25C7	184CD1	43.68
190E1	25C7	20N	74.85	190E1	25C7	20CA	91.61
210E1	25C7	21N	49.78	210E1	25C7	20N	86.97
210E1	25C7	21NE2	27.10	210E1	25C7	20CA	70.13
21N	25C7	20N	44.95	21N	25C7	21NE2	38.79
21N	25C7	20CA	28.96	184CD1	25C7	20N	77.27
184CD1	25C7	20CA	92.17	20N	25C7	21NE2	62.89
20N	25C7	20CA	17.26	21 <b>NE2</b>	25C7	20CA	47.61
200	2508	184NE1	98.56	200	2508	19NE2	64.94
200	2508	19CD	64.80	200	2508	19CG	49.58
200	2508	184CD1	86.05	200	2508	190E1	77.25
184NE1	2508	19NE2	65.95	184NE1	2508	19CD	51.39
184NE1	2508	19CG	55.21	184NE1	2508	184CE2	15.43
184NE1	2508	184CD1	14.81	184NE1	2508	190E1	39.31
19NE2	2508	19CD	16.73	19NE2	2508	19CG	29.82
19NE2	2508	184CE2	79.52	19NE2	2508	184CD1	66.86
19NE2	2508	190E1	26.64	1.9CD	2508	19CG	18.32
19CD	2508	184CE2	66.04	19CD	2508	184CD1	50.50

				BLE XVIII			
19CD	2508	190E1	13.90	19 <b>C</b> G	2508	184CE2	70.62
19CG	2508	184CD1	48.50	19CG	2508	190E1	28.11
184CE2	2508	184CD1	26.04	184CE2	2508	190E1	53.16
184CD1	2508	190E1	41.64	184NE1	25C9	19NE2	75.95
184NE1	25C9	19CD	58.34	184NE1	25C9	184CE2	18.51
184NE1	25C9	184CZ2	36.32	184NE1	25C9	190E1	46.11
184NE1	25C9	162CE1	47.88	184NE1	25C9	184CD1	13.15
184NE1	25C9	19CG	58.03	19NE2	25C9	19CD	18.37
19NE2	25C9	184CE2	93.09	19NE2	25C9	190E1	30.69
19NE2	25C9	162CE1	65.23	19NE2	25 <b>C</b> 9	184CD1	73.02
19NE2	25C9	19CG	30.00	19CD	25C9	184CE2	76.16
19CD	25C9	184CZ2	89.88	19CD	25C9	190E1	16.50
19CD	25C9	162CE1	58.87	19CD	25C9	184CD1	54.66
19CD	25C9	19CG	17.52	184CE2	25C9	184CZ2	19.20
184CE2	25C9	190E1	62.52	184CE2	25C9	162CE1	49.95
184CE2	25C9	184CD1	28.40	184CE2	25C9	19CG	76.51
184CZ2	25C9	190E1	74.32	184CZ2	25C9	162CE1	47.31
184CZ2	25C9	184CD1	47.47	184CZ2	25C9	19CG	93.64
190E1	25C9	162CE1	43.44	190E1	25C9	184CD1	46.14
190E1	25C9	19CG	29.61	162CE1	25C9	184CD1	58.96
162CE1	25 <b>C</b> 9	19CG	72.86	184CD1	25C9	19CG	50.05
184NE1	25010	184CE2	25.08	184NE1	25010	184CZ2	48.91
184NE1	25010	190E1	62.37	184NE1	25010	19CD	75.84
184NE1	25010	184CD1	15.34	184NE1	25010	162CE1	66.14
184NE1	25010	19NE2	95.76	184NE1	25010	162ND1	81.12
184NE1	25010	19CG	70.87	184NE1	25010	184CD2	22.23
184NE1	25010	162NE2	52.65	184NE1	25010	184CH2	53.72
184NE1	25010	184CG	14.63	184CE2	25010	184CZ2	25.03
184CE2	25010	190E1	84.37	184CE2	25010	184CD1	37.35
184CE2	25010	162CE1	66.88	184CE2	25010	162ND1	77.00
184CE2	25010	19CG	95.89	184CE2	25010	184CD2	10.58
184CE2	25010	162NE2	51.22	184CE2	25010	184CH2	28.91
184CE2	25010	184CG	26.65	184CZ2	25010	190E1	99.91
184CZ2		184CD1	62.29	184CZ2		162CE1	61.92
184CZ2	25010		65.32	184CZ2		184CD2	33.91
184CZ2		162NE2	48.17	184CZ2		184CH2	7.17
184CZ2		184CG	51.50	190E1	25010	19CD	19.97
190E1		184CD1	<b>59.9</b> 0	190E1		162CE1	56.15
190E1	25010	19NE2	35.40	190E1		162ND1	68.84
190E1	25010	19CG	33.29	190E1		184CD2	84.51
190E1		162NE2	60.20	190E1		184CG	72.32
19CD		184CD1	68.82	19CD		162CE1	74.19
19CD	25010	19NE2	20.25	1.9CD		162ND1	84.51
19CD	25010	19CG	19.19	1900	25010	184CD2	97.71

			TAI	BLE XVIII			
19CD	25010	162NE2	79.91	19CD	25010	184CG	82.24
184CD1	25010	162CE1	78.10	184CD1	25010	19NE2	89.03
184CD1	25010	162ND1	94.07	184CD1	25010	19CG	59.77
184CD1	25010	184CD2	30.63	184CD1	25010	162NE2	66.09
184CD1	25010	184CH2	66.12	184CD1	25010	184CG	13.45
162CE1	25010	19NE2	78.81	162CE1	25010	162ND1	16.74
162CE1	25010	19CG	89.36	162CE1	25010	184CD2	75.74
162CE1	25010	162NE2	15.66	162CE1	25010	184CH2	68.07
162CE1	25010	184CG	80.42	19NE2	25010	162ND1	83.89
19NE2	25010	19CG	33.67	19NE2	25010	162NE2	89.03
162ND1	25010	184CD2	86.96	162ND1	25010	162NE2	28.50
162ND1	25010	184CH2	70.03	162ND1	25010	184CG	94.80
19CG	25010	184CD2	90.39	19CG	25010	162NE2	91.46
19CG	25010	184CG	72.81	184CD2	25010	162NE2	60.21
184CD2	25010	184CH2	36.32	184CD2	25010	184CG	18.11
162NE2	25010	184CH2	54.84	162NE2	25010	184CG	66.33
184CH2	25010	184CG	54.39	162ND1	25C11	162CE1	18.33
162ND1	25C11	184CZ2	57.54	162ND1	25C11	19NE2	74.67
162ND1	25C11	162CG	12.31	162CE1	25C11	184CZ2	47.54
162CE1	25C11	19NE2	62.12	162CE1	25C11	162CG	26.24
184CZ2	25C11	19NE2	86.59	184CZ2	25C11	162CG	53.23
19NE2	25C11	162CG	86.37	162ND1	25C12	1610	52.69
1610D1	25C13	162ND1	77.53	1610D1	25C13	161CG	13.50
1610D1	25C13	1610	49.60	1610D1	25C13	162CB	47.00
1610D1	25C13	162CG	64.44	1610D1	25C13	1370	63.21
162ND1	25C13	161CG	86.85	162ND1	25C13	1610	55.29
162ND1	25C13	184CZ2	52.94	162ND1	25C13	162CB	31.13
162ND1	25C13	162CG	16.56	162ND1	25C13	1370	91.79
161CG	25C13	1610	48.18	161CG	25C13	162CB	57.82
161CG	25C13	162CG	75.53	161CG	25C13	1370	72.06
1610	25C13	162CB	45.12	1610	25C13	162CG	55.63
184CZ2	25C13	162CB	67.90	184CZ2	25C13	162CG	52.88
184CZ2	25C13	1370	65.96	162CB	25C13	162CG	17.71
162CB		1370	73.09	162CG	25C13		76.90
1610D1		162CB	67.52	1610D1		162CG	91.93
1610D1	25C14		65.79	1610D1	25C14		49.33
1610D1		162CA	65.70	1610 <b>D1</b>		161CG	13.81
1610D1	25C14		48.26	1610D1		137CB	61.84
1610D1		161CB	32.59	1610D1		162CD2	97.90
151001	25C14		54.56	1610D1		161CA	38.35
1610D1	25C14		68.99	1610D1		137CA	49.42
162ND1		162CB	44.41	162ND1		162CG	23.28
162ND1	25C14		75.37	162ND1	25C14		79.50
162ND1	25C14	162CA	48.35	162ND1	25C14	1.62N	68.54

			TAE	LE XVIII			
162ND1	25C14	162CE1	14.58	162ND1	25C14	137CB	81.24
162ND1	25C14	162CD2	28.11	162ND1	25C14	184CZ2	60.58
162ND1	25C14	162NE2	21.94	162ND1	25C14	161CA	95.48
162ND1	25C14	137CA	97.09	162CB	25C14	162CG	25.05
162CB	25C14	1610	62.89	162CB	25C14	161C	53.94
162CB	25C14	162CA	21.95	162CB	25C14	161CG	76.82
162CB	25C14	162N	35.96	162CB	25C14	162CE1	55.50
162CB	25C14	137CB	53.40	162CB	25C14	161CB	80.35
162CB	25C14	162CD2	35.20	162CB	25C14	184CZ2	83.43
162CB	25C14	137C	81.05	162CB	25C14	162NE2	49.18
162CB	25C14	161CA	65.39	162CB	25C14	1370	88.14
162CB	25C14	137CA	63.27	162CG	25C14	1610	77.15
162CG	25C14	161C	73.93	162CG	25C14	162CA	38.93
162CG	25C14	162N	57.98	162CG	25C14	162CE1	31.20
162CG	25C14	137CB	59.50	162CG	25C14	162CD2	13.71
162CG	25C14	184CZ2	62.16	162CG	25C14	137C	90.53
162CG	25C14	162NE2	24.37	162CG		161CA	87.82
162CG	25C14	1370	91.83	162CG	25C14	137CA	74.49
1610	25C14	161C	17.83	1610	25C14	162CA	41.16
1610	25C14	161CG	59.81	1610	25C14	162N	32.33
1610	25C14	162CE1	89.00	1610		161CB	43.28
1610	25C14	162CD2	90.86	1610	25C14	162NE2	95.79
1610	25C14	161CA	28.30	161C	25C14	162CA	35.01
161C	25C14	161CG	45.86	161C	25C14	162N	18.49
161C	25C14	162CE1	94.05	161C	25C14		91.74
161C	25C14	161CB	34.16	161C	25C14	162CD2	87.05
161C	25C14	162NE2	96.49	161C		161CA	16.14
161C	25C14	137CA	88.79	162CA	25C14	161CG	70.29
162CA	25C14	162N	20.32	162CA	25C14		62.52
162CA	25C14	137CB	73.16	162CA	25C14		66.66
1.62CA	25C14	162CD2	52.11	162CA	25C14	137C	96.91
162CA	25C14	162NE2		162CA	25C14	161CA	49.32
162CA	25C14		79.58	161CG	25C14	162N	50.67
161CG		137CB	75.47	161CG		161CB	20.44
161CG	25C14	137C	64.73	161CG	25C14		31.54
161CG		1370	78.60	161CG		137CA	62.23
162N	25C14	162CE1	82.81	162N		137CB	75.61
1.62N	25C14	161CB	46.58	162N		162CD2	70.31
162N		137C	90.94	162N		162NE2	81.86
162N	25C14	161CA	23.85	162N		137CA	76.03
162CE1		137CB	81.53	162CE1		162CD2	28.99
162CE1		184CZ2	47.31	162CE1		162NE2	14.94
162CE1		137CA	98.37	137CB		161CB	93.78
137CB	25C14	162CD2	53.66	137CB	25C14	184CZ2	66.27

			TAI	BLE XVIII			
137CB	25C14	137C	31.04	137CB	25C14	162NE2	66.78
137CB	25C14	161CA	91.93	137CB	25C14	1370	34.84
137CB	25C14	137CA	16.94	161CB	25C14	137C	85.13
161CB	25C14	161CA	18.06	161CB	25C14	1370	98.82
161CB	25C14	137CA	82.01	162CD2	25C14	184CZ2	49.12
162CD2	25C14	137C	84.06	162CD2	25C14	162NE2	16.33
162CD2	25C14	1370	82.53	162CD2	25C14	137CA	70.12
184CZ2	25C14	137C	79.92	184CZ2	25C14	162NE2	40.56
184CZ2	25C14	1370	67.98	184CZ2	25C14	137CA	79.44
137C	25C14	162NE2	95.36	137C	25C14	161CA	92.71
137C	25C14	1370	14.52	137C	25C14	137CA	17.79
162NE2	25C14	1370	90.59	162NE2	25C14	137CA	83.69
161CA	25C14	137CA	84.51	1370	25C14	137CA	28.01
1370	25C15	184CZ2	82.18	1370	25C15	184CH2	65.00
1370	25C15	137C	15.27	1370	25C15	1610D1	65.35
1370	25C15	138CA	35.59	1370	25C15	138N	27.26
1370	25C15	137CB	35.12	184CZ2	25C15	184CH2	18.93
184CZ2	25C15	137C	89.79	184CZ2	25C15	137CB	65.87
184CH2	25C15	137C	74.80	184CH2	25C15	138N	90.34
184CH2	25C15	137CB	56.59	137C	25C15	1610D1	50.24
137C	25C15	138CA	30.23	137C	25C15	138N	15.68
137C	25C15	137CB	30.32	1610D1	25C15	138CA	56.67
1610D1	25C15	138N	45.35	1610D1	25C15	137CB	48.96
138CA	25C15	138N	17.26	138CA	25C15	137CB	59.96
138N	25C15	137CB	43.35	162ND1	25C16	25 <b>S</b> G	47.96
162ND1	25C16	19NE2	81.02	162ND1	25C16	162CE1	17.42
162ND1	25C16	1610	59.00	25SG	25C16	19NE2	64.48
25 <i>S</i> G	25C16	162CE1	51.01	25SG	25C16	23CA	79.30
25SG	25C16	1610	61.60	19NE2	25C16	162CE1	65.94
19NE2	25C16	23CA	51.63	162CE1	25C16	1610	76.23
23CA	25017	19NE2	63.28	23CA	25017	23C	18.00
23CA	25017	23N	16.04	23CA	25017	220	36.68
23CA	25017	25SG	86.81	23CA	25017	22C	27.89
23CA	25017	19CD	75.23	23CA	25017	230	26.79
23CA	25017	24N	26.12	19NE2	25017	23C	56.89
19NE2	25017	23N	63.52	19NE2	25017	220	37.86
19NE2	25017	25SG	54.03	19NE2	25017	22C	51.98
19NE2	25017	19CD	12.01	19NE2	25017	230	67.69
19NE2	25017	24N	42.39	23C	25017	23N	32.96
23C	25017	220	43.34	23C	25017	25SG	68.82
23C	25017	22C	40.37	23C	25017	19CD	63.57
23C	25017	230	13.76	23C	25017	24N	14.50
23N	25017	220	28.32	23N	25017	22C	15.59
23N	25017	19CD	74.62	23N	25017	230	42.84

			TAE	LE XVIII			
23N	25017	24N	36.39	220	25017	25SG	92.75
220	25017	22C	14.54	220	25017	19CD	47.50
220	25017	230	57.03	220	25017	24N	35.11
25SG	25017	19CD	63.67	25SG	25017	230	64.10
25 <i>S</i> G	25017	24N	64.69	22C	25017	19CD	61.99
22C	25017	230	52.87	22C	25017	24N	37.51
19CD	25017	230	78.75	19CD	25017	24N	54.07
230	25017	24N	26.33	25 <i>S</i> G	25N18	162ND1	59.84
25 <i>S</i> G	25N18	1610	84.84	25SG	25N18	162CE1	58.94
25 <i>S</i> G	25N18	25CB	19.83	25SG	25N18	162CG	65.68
25 <i>S</i> G	25N18	161C	84.84	25SG	25N18	162CA	63.77
25SG	25N18	162CB	72.00	25SG	25N18	19NE2	64.56
162ND1	25N18	1610	77.72	162ND1	25N18	162CE1	17.48
162ND1	25N18	25CB	52.80	162ND1	25N18	162CG	12.09
162ND1	25N18	161C	70.93	162ND1	25N18	162CA	42.08
162ND1	25N18	162CB	29.83	162ND1	25N18	19NE2	80.21
1610	25N18	162CE1	95.07	1610	25N18	162CG	66.81
1610	25N18	161C	7.83	1610	25N18	162CA	36.26
1610	25N18	162CB	49.33	162CE1	25N18	25CB	45.72
162CE1	25N18	162CG	29.21	162CE1	25N18	161C	88.39
162CE1	25N18	162CA	59.10	162CE1	25N18	162CB	47.16
162CE1	25N18	19NE2	63.44	25CB	25N18	162CG	62.19
25CB	25N18	161C	98.68	25CB	25N18	162CA	72.18
25CB	25N18	162CB	74.19	25CB	25N18	19NE2	48.00
162CG	25N18	161C	59.70	162CG	25N18	162CA	32.47
162CG	25N18	162CB	17.97	162CG	25N18	19NE2	92.29
161C	25N18	162CA	30.47	161C	25N18	162CB	41.99
162CA	25N18	162CB	18.49	25SG	25C19	1610	87.86
25 <i>S</i> G	25C19	25CB	17.43	25 <b>SG</b>	25C19	162ND1	51.44
25 <b>S</b> G	25C19	230	85.01	25 <b>SG</b>	25C19	23C	84.49
25SG	25C19	25N	40.23	25 <b>S</b> G	25C19	161C	84.17
25SG	25C19	162CE1	48.33	25 <b>S</b> G	25C19	19NE2	65.89
1610	·25C19	162ND1	65.41	1610	25C19	161C	3.96
1610		162CE1	79.95	25CB		162ND1	50.53
25CB	25C19	23CA	87.26	25CB	25C19	230	79.49
25CB	25C19	23C	74.66	25CB	25C19	25N	31.30
25CB	25C19		97.21	25 <b>CB</b>	25C19		42.12
25CB	25C19	19NE2	48.60	162ND1	25C19	25N	81.01
162ND1	25C19		61.89	162ND1		162CE1	14.54
162ND1	25C19	19NE2	72.57	23 <b>CA</b>	25C19	230	31.62
23CA	25C19	23C	19.66	23CA	25C19	25N	60.25
23CA	25C19	19NE2	49.27	230	25C19	23C	15.91
230	25C19	25N	48.22	230	25C19		64.33
23C	25C19	25N	44.48	23C	25C19	19NE2	49.75

			TAI	BLE XVIII			
25N	25C19	162CE1	70.28	25N	25C19	19NE2	41.94
161C	25C19	162CE1	76.42	162CE1	25C19	19NE2	58.03
19NE2	25 <b>N2</b> 0	184NE1	61.55	19NE2	25N20	184CZ2	88.39
19NE2	25N20	162CE1	59.51	19NE2	25N20	19CD	15.18
184NE1	25 <b>N2</b> 0	184CZ2	30.44	184NE1	25N20	162CE1	42.35
184NE1	25N20	19CD	46.54	184CZ2	25N20	162CE1	43.21
184CZ2	25N20	19CD	74.46	162CE1	25N20	19CD	52.22
1610	25C21	25 <b>S</b> G	96.09	1610	25C21	161C	6.24
1610	25C21	162CA	33.10	1610	25C21	162N	17.36
1610	25C21	162ND1	61.84	1610	25C21	25CB	99.78
1610	25C21	163N	59.11	25SG	25C21	161C	94.55
25 <i>S</i> G	25C21	162CA	63.31	25SG	25C21	162N	80.59
25SG	25C21	162ND1	43.09	25SG	25C21	25CB	6.39
25SG	25C21	163N	44.50	161C	25C21	162CA	31.26
161C	25C21	162N	14.07	161C	25C21	162ND1	63.41
161C	25C21	25CB	98.77	161C	25C21	163N	55.24
162CA	25C21	162N	17.54	162CA	25C21	162ND1	37.75
162CA	25C21	25CB	67.54	162CA	25C21	163N	28.93
162N	25C21	162ND1	52.75	162N	25C21	25CB	85.01
162N	25C21	163N	41.75	162ND1	25C21	25CB	43.42
162ND1	25C21	163N	45.75	25CB	25C21	163N	50.47
25 <i>S</i> G	25C22	25CB	32.60	25 <b>S</b> G	25C22	25N	72.57
25 <i>S</i> G	25C22	25CA	49.33	25 <b>S</b> G	25C22	19NE2	93.81
25 <b>S</b> G	25C22	26N	55.18	25 <i>S</i> G	25C22	162ND1	47.43
25 <i>S</i> G	25C22	24C	82.40	25SG	25C22	25C	44.53
25 <b>S</b> G	25C22	162CE1	48.15	25SG	25C22	1610	75.53
25SG	25C22	26CD1	94.13	25 <b>S</b> G	25C22	190E1	68.36
25 <b>\$</b> G	25C22	19CD	81.56	25CB	25C22	25N	46.75
25CB	25C22	25CA	24.63	25CB	25C22	19NE2	63.05
25CB	25C22	24N	85.73	25CB	25C22	26N	52.12
25CB	25C22	162ND1	56.93	25CB	25C22	24C	57.01
25CB	25C22	25C	34.57	25CB	25C22	162CE1	46.25
25CB	25C22	24CA	75.71	25CB	25C22	26CD1	92.36
25CB	25C22	190E1	41.24	25CB	25C22	19CD	51.95
25 <b>N</b>	25C22	25CA	23.55	25 <b>N</b>	25C22	230	65.42
25 <b>N</b>	25C22	23C	60.13	2 <b>5N</b>	25C22	23CA	77.28
25N	25C22	19NE2	54.87	25N	25C22	24N	43.20
25 <b>N</b>	25C22	26N	39.45	25 <b>N</b>	25C22	24C	10.31
25N	25C22	25C	33.15	25N	25C22	162CE1	88.28
25 <b>N</b>	25C22	24CA	29.39	25N	25C22	26CD1	58.€4
25N	25C22	190E1	57.16	25N	25C22	19CD	54.20
25CA	25C22	230	88.34	25 <b>CA</b>	25C22	23C	83.54
25CA	25C22	23CA	99.22	25CA	25C22	19NE2	61.73
25CA	25C22	24N	66.05	25 <b>CA</b>	25C22	26N	35.11

			TAB	LE XVIII			
25CA	25C22	162ND1	81.55	25CA	25C22	24C	33.75
25CA	25C22	25C	19.23	25CA	25C22	162CE1	69.99
25CA	25C22	24CA	52.91	25CA	25C22	26CD1	70.42
25CA	25C22	190E1	50.96	25CA	25C22	19CD	55.14
230	25C22	23C	19.26	230	25C22	23CA	35.89
230	25C22	19NE2	79.72	230	25C22	24N	32.92
230	25C22	26N	78.15	230	25C22	24C	55.15
230	25C22	25C	88.29	230	25C22	24CA	36.77
230	25C22	26CD1	45.35	230	25C22	19CD	90.80
23C	25C22	23CA	22.09	23C	25C22	19NE2	61.06
23C	25C22	24N	18.52	23C	25C22	26N	84.89
23C	25C22	24C	50.29	23C	25C22	25C	89.80
23C	25C22	24CA	31.49	23C	25C22	26CD1	60.86
23C	25C22	190E1	86.44	23C	25C22	19CD	72.74
23CA	25C22	19NE2	57.32	23CA	25C22	24N	34.39
23CA	25C22	24C	68.55	23CA	25C22	24CA	51.33
23CA	25C22	26CD1	81.00	23CA	25C22	190E1	84.42
23CA	25C22	19CD	70.13	19NE2	25C22	24N	48.45
19NE2	25C22	26N	93.13	19NE2	25C22	162ND1	81.54
19NE2	25C22	24C	55.55	19NE2	25 <b>C</b> 22	25C	80.63
19NE2	25C22	162CE1	65.93	19NE2	25C22	24CA	57.12
19NE2	25C22	190E1	27.13	19NE2	25C22	19CD	12.81
24N	25C22	26N	74.25	24N	25C22	24C	34.17
24N	25C22	25C	75.06	24N	25C22	24CA	17.99
24N	25C22	26CD1	62.40	24N	25C22	190E1	70.97
24N	25C22	19CD	58.41	2 <b>6N</b>	25C22	24C	43.27
26N	25C22	25C	17.72	26N	25C22	162CE1	96.42
26N	25C22	24CA	56.30	26N	25C22	26CD1	41.56
26N	25C22	190E1	86.07	26N	25C22	19CD	89.17
162ND1	25C22	25C	87.29	162ND1	25C22	162CE1	16.50
162ND1	25C22	1610	57.76	162ND1	25C22	190E1	57.61
162ND1	25C22	19CD	70.43	24C	25C22	25C	40.95
24C	25C22	162CE1	97.34	24C	25C22	24CA	19.17
24C	25C22	26CD1	53.28	24C	25C22	190E1	63.13
24C	25C22	19CD	57.60	25C	25C22	162CE1	79.88
25C	25C22	24CA	58.50	25C	25C22	26CD1	58.12
25C	25C22	190E1	69.35	25C	25C22	19CD	74.36
162CE1	25C22		74.20	162CE1	25C22		41.17
162CE1	25C22	19CD	54.32	24CA	25C22		49.39
24CA	25C22	190E1	73.34	24CA	25C22		63.76
190E1	25C22	19CD	14.35	25SG	25023		75.69
25 <b>S</b> G	25023	25CB	37.11	25\$G	25023		54.99
25SG	25023	24C	90.69	25SG	25023		83.77
25SG	25023	25C	48.15	25SG	25023	26N	51.79

			TAB	LE XVIII			
25 <b>S</b> G	25023	162ND1	44.43	25SG	25023	162CE1	52.19
25SG	25023	240	85.50	25N	25023	25CB	51.43
25N	25023	19NE2	75.55	25N	25023	23C	76.45
25N	25023	24N	57.15	25N	25023	230	76.49
25N	25023	25CA	24.40	25N	25023	24C	15.43
25N	25023	24CA	38.06	25N	25023	19CD	72.15
25N	25023	190E1	71.82	25N	25023	220	86.12
25N	25023	25C	27.77	25N	25023	26N	33.20
25N	25023	23N	97.97	25N	25023	162CE1	94.44
25N	25023	240	9.85	25N	25023	22C	91.69
25CB	25023	19NE2	80.56	25CB	25023	25CA	27.06
25CB	25023	24C	66.23	25CB	25023	24CA	89.22
25CB	25023	19CD	67.04	25CB	25023	190E1	52.55
25CB	25023	25C	33.18	25CB	25023	26N	48.61
25CB	25023	162ND1	52.03	25CB	25023	162CE1	45.88
25CB	25023	240	59.52	19NE2	25023	23C	84.54
19NE2	25023	24N	66.47	19NE2	25023	23CA	78.32
19NE2	25023	25CA	77.61	19NE2	25023	24C	71.54
19NE2	25023	24CA	74.20	19NE2	25023	19CD	14.67
19NE2	25023	190E1	32.03	19NE2	25023	220	39.73
19NE2	25023	25C	93.21	19NE2	25023	23N	68.79
19NE2	25023	162ND1	89.64	19NE2	25023	162CE1	73.89
19NE2	25023	240	69.92	19NE2	25023	22C	53.32
23C	25023	24N	24.58	23C	25023	23CA	28.13
23C	25023	230	21.76	23C	25023	24C	61.59
23C	25023	24CA	38.61	23C	25023	19CD	97.49
23C	25023	220	49.19	23C	25023	25C	99.27
23C	25023	26N	90.14	23C	25023	23N	30.80
23C	25023	240	68.35	23C	25023	22C	39.19
24N	25023	23CA	44.35	24N	25023	230	40.46
24N	25023	25CA	80.66	24N	25023	24C	41.72
24N	25023	24CA	20.87	24N	25023	19CD	77.11
24N	25023	190E1	91.57	24N	25023	220	41.76
24N	25023	25C	83.71	24N	25023	26N	80.39
24N	25023	23N	41.16	24N	25023	240	47.80
24N	25023	22C	39.24	23CA	25023	230	43.14
23CA	25023	24C	85.93	23CA	25023	24CA	63.70
23CA	25023	19CD	92.98	23 <b>CA</b>	25023	220	38.70
23CA	25023	23N	9.54	23CA	25023	240	92.15
23CA	25023	22C	25.05	230	25023	25CA	99.19
230	25023	24C	64.48	230	25023	24CA	44.29
230	25023	220	70.90	230	25023	25C	91.92
230	25023	26N	78.46	230	25023	23N	49.19
230	25023	240	71.18	230	25023	22C	60.24

			TAE	BLE XVIII			
25CA	25023	24C	39.48	25CA	25023	24CA	62.38
25CA	25023	19CD	68.41	25CA	25023	190E1	60.76
25CA	25023	25C	15.95	25CA	25023	26N	32.35
25CA	25023	162ND1	79.08	25CA	25023	162CE1	71.50
25CA	25023	240	32.98	24C	25023	24CA	22.98
24C	25023	19CD	72.19	24C	25023	190E1	76.64
24C	25023	220	73.41	24C	25023	25C	42.54
24C	25023	26N	44.02	24C	25023	23N	82.66
24C	25023	240	6.88	24C	25023	22C	77.34
24CA	25023	19CD	80.54	24CA	25023	190E1	90.57
24CA	25023	220	60.20	24CA	25023	25C	63.31
24CA	25023	26N	59.62	24CA	25023	23N	61.69
24CA	25023	240	29.75	24CA	25023	22C	59.80
19CD	25023	190E1	17.43	19CD	25023	220	54.37
19CD	25023	25C	84.36	19CD	25023	26N	99.96
19CD	25023	23N	83.45	19CD	25023	162ND1	76.89
19CD	25023	162CE1	60.68	19CD	25023	240	68.84
19CD	25023	22C	67.98	190E1	25023	220	71.76
190E1	25023	25C	76.10	190E1	25023	26N	93.05
190E1	25023	162ND1	60.98	190E1	25023	162CE1	44.53
190E1	25023	240	71.61	190E1	25023	22C	85.33
220	25023	23N	29.15	220	25023	240	76.62
220	25023	22C	13.64	25C	25023	26N	17.16
25C	25023		82.30	25C	25023	162CE1	78.86
25C	25023	240	37.62	26N	25023	162ND1	92.78
26N	25023	162CE1	92.75	26N	25023	240	41.67
23N	25023	240	88.33	23N	25023	22C	15.51
162ND1	25023	162CE1	16.46	240	25023	22C	81.85
65CA	25C24	6 <b>6N</b>	33.46	65CA	25C24	65C	20.36
65CA	25C24	540	36.58	65CA	25C24	660	68.87
66N	25C24	65C	17.64	6 6 N	25C24	640	67.62
66N	25C24	660	36.45	65C	25C24	640	50.58
65C	25C24	660	53.76	660	25C25	6 <b>6N</b>	40.06
660	25C25	66C	8.90	660	25C25	65CA	70.27
660	25C25	65C	53.58	1610	25C25	161C	14.86
1610	25C25		28.19	1610	25C25	25 <b>S</b> G	60.39
66N	25C25	6 <b>6C</b>	31.52	6 <b>SN</b>	25C25	65CA	30.22
66N	25C25	25 <b>S</b> G	91.38	6 <b>6N</b>	25C25	65C	14.24
161C		161CA	1.7.63	161C	25C25	25 <i>S</i> G	69.22
66C	25C25	65CA	61.71	<b>୬</b> ୧୯	25C25	65C	44.80
65CA	25C25	25 <i>S</i> G	82.36	65CA	25C25	65C	17.73
161CA	25C25	25 <b>S</b> G	86.68	25 <b>S</b> G	25C25	65C	92.33
660	25C26	66C	3.30	660	25C26		37.58
1610	25C26	161C	1584	1610	25C26	163N	56.85

			TAI	BLE XVIII			
1610	25C26	1600	61.70	1610	25C26	163CB	84.35
1610	25C26	162N	27.01	1610	25C26	161CA	28.97
161C	25C26	163N	55.26	161C	25C26	1600	50.24
161C	25C26	163CB	84.79	161C	25C26	162N	15.32
161C	25C26	161CA	17.79	163N	25C26	1600	95.75
163N	25C26	163CB	29.64	163N	25C26	162N	42.53
163N	25C26	161CA	71.03	66C	25C26	163CB	89.72
1600	25C26	162N	56.11	1600	25C26	161CA	33.06
163CB	25C26	162N	72.06	162N	25C26	161CA	28.56
1600	25C27	160C	16.28	1600	25C27	161CA	39.86
1600	25C27	161C	59.69	1600	25C27	161N	29.94
1600	25C27	1610	70.88	1600	25C27	160CB	39.29
1600	25C27	134CB	91.77	1600	25C27	162N	65.04
160C	25C27	161CA	33.56	160C	25C27	161C	50.71
160C	25C27	161N	17.29	160C	25C27	1610	64.68
160C	25C27	160CB	32.70	160C	25C27	134CB	77.93
160C	25C27	162N	52.23	161CA	25C27	161C	20.25
161CA	25C27	161N	19.14	161CA	25C27	1610	31.54
161CA	25C27	160CB	64.50	161CA	25C27	134CB	90.77
161CA	25C27	162N	30.47	161C	25C27	161N	33.69
161C	25C27	1610	15.99	161C	25C27	160CB	77.36
161C	25C27	134CB	85.39	161C	25C27	162N	15.75
161N	25C27	1610	48.38	161N	25C27	160CB	45.58
161N	25C27	134CB	77.13	161N	25C27	162N	35.42
1610	25C27	160CB	93.10	1610	25C27	134CB	96.47
1610	25C27	162N	27.78	660	25C27	209CD2	83.62
160CB	25C27	134CB	56.56	160CB	25C27	162N	71.89
160CB	25C27	209CD2	69.27	134CB	25C27	162N	69.95
134CB	25C27	209CD2	49.25	1600	25C28	160C	11.12
1600	25C28	161CA	34.92	1600	25C28	161N	22.64
160C	25C28	161CA	30.68	160C	25C28	161N	14.74
67CE1	25C28	660	57.93	67CE1	25C28	67CD1	16.20
161CA ·	25C28	161N	17.22	660	25C28	67CD1	44.47
209CD2	25C29		68.69	209CD2	25C29	160CB	90.30
209CD2	25C29	67CD1	65.58	209CD2	25C29	6 <b>6</b> 0	98.85
209CD2	25C29	209CG	7.93	209CD2	25C29	67CE1	70.23
1.34CB	25C29	160CB	68.29	134CB	25C29	160C	85.31
134CB	25C29	209CG	62.21	1500	25C29	160CB	42.09
1600	25C29	160C	15.52	160CB	25C29	160C	33.72
160CB	25C29		83.59	67CD1	25C29	660	45.57
67CD1	25C29	209CG	73.48	67CD1	25C29	67CE1	16.51
660	25C29	67CE1	56.12	209CG	25C29	67CE1	77.57
660	25C30	6 GN	46.15	6 <b>60</b>	25C30	65CA	82.36
660	25C30	26CD1	69.05	660	25C30	26CB	54.83

TABLE XVIII									
660	25C30	65C	61.73	660	25C30	66C	12.32		
660	25C30	66CA	31.12	660	25C30	26CG	57.20		
660	25C30	26N	86.09	660	25C30	163CB	86.18		
66N	25C30	65CA	36.68	66N	25C30	26CD1	45.40		
66N	25C30	26CB	62.49	66N	25C30	65C	16.57		
66N	25C30	66C	34.22	66N	25C30	66CA	15.04		
66N	25C30	26CG	48.20	66N	25C30	26N	85.31		
25SG	25C30	26CD1	80.34	25 <i>S</i> G	25C30	26CB	78.23		
25 <i>S</i> G	25C30	1610	67.14	25SG	25C30	26CG	83.17		
25 <b>S</b> G	25C30	26N	47.19	25SG	25C30	163CB	52.42		
25SG	25C30	163N	44.84	65CA	25C30	26CD1	53.97		
65CA	25C30	26CB	86.07	65CA	25C30	65C	20.64		
65CA	25C30	66C	70.78	65CA	25C30	66CA	51.58		
65CA	25C30	26CG	67.39	65CA	25C30	26N	93.38		
26CD1	25C30	26CB	34.81	26CD1	25C30	65C	49.80		
26CD1	25C30	66C	59.13	26CD1	25C30	66CA	50.39		
26CD1	25C30	26CG	17.00	26CD1	25C30	26N	42.15		
26CD1	25C30	163CB	83.57	26CB	25C30	65C	74.87		
26CB	25C30	66C	51.73	26CB	25C30	66CA	56.97		
26CB	25C30	26CG	18.84	26CB	25C30	26N	31.41		
26CB	25C30	163CB	54.44	26CB	25C30	163N	82.52		
65C	25C30	66C	50.25	65C	25C30	66CA	31.10		
65C	25C30	26CG	58.02	65C	25C30	26N	91.94		
1610	25C30	163CB	83.35	1610	25C30	163N	54.69		
66C	25C30	66CA	19.22	66C	25C30	26CG	49.60		
66C	25C30	26N	82.89	66C	25C30	163CB	92.24		
66CA	25C30	26CG	47.25	66CA	25C30	26N	84.74		
26CG	25C30	26N	37.73	26CG	25C30	163CB	71.86		
26CG	25C30	163N	98.68	26N	25C30	163CB	44.77		
26N	25C30	163N	65.10	163CB	25C30	163N	28.72		
660	25031	66N	61.37	660	25031	26CB	77.21		
660	25031	26CD1	95.50	660	25031	66C	18.26		
660	25031	26CG	80.22	660	25031	66CA	41.13		
660	25031	65C	76.22	660	25031	65CA	98.52		
660	25031	26CA	93.42	660	25031	26NE1	92.30		
660	25031	67N	17.71	660	25031	26CD2	73.43		
660	25031	650	71.00	66N	25031	26CB	86.34		
66N	25031	26CD1	59.28	66N	25031	66C	45.73		
66N	25031	26CG	65.50	66N	25031	66CA	21.43		
66N	25031	65C	1.5.23	6 <b>6N</b>	25031	65CA	39.07		
66N	25031	26NE1	49.90	66N	25031	67N	52.44		
66N	25031	26CD2	57.23	66N	25031	650	11.55		
26CB	25031	26CD1	45.93	26CB	25031	66C	70.12		
26CB	25031	26CG	25.17	2.6CB	25031	66CA	76.25		

			TAE	BLE XVIII			
26CB	25031	65C	96.29	26CB	25031	26N	36.99
26CB	25031	26CA	18.87	26CB	25031	25 <b>S</b> G	86.19
26CB	25031	26NE1	53.75	26CB	25031	67N	63.56
26CB	25031	163CB	61.81	26CB	25031	26CD2	30.74
26CB	25031	650	94.86	26CD1	25031	66C	78.62
26CD1	25031	26CG	22.75	26CD1	25031	66CA	65.15
26CD1	25031	65C	60.18	26CD1	25031	65CA	62.48
26CD1	25031	26N	50.53	26CD1	25031	26CA	52.01
26CD1	25031	25SG	86.23	26CD1	25031	26NE1	10.06
26CD1	25031	67N	77.85	26CD1	25031	163CB	97.88
26CD1	25031	26CD2	24.53	26CD1	25031	650	61.57
66C	25031	26CG	66.43	66C	25031	66CA	24.43
66C	25031	65C	61.60	66C	25031	65CA	84.54
66C	25031	26CA	88.49	66C	25031	26NE1	74.51
66C	25031	67N	8.31	66C	25031	26CD2	58.54
66C	25031	650	56.49	26CG	25031	66CA	61.97
26CG	25031	65C	72.65	26CG	25031	65CA	81.58
26CG	25031	26N	45.87	26CG	25031	26CA	37.48
26CG	25031	25 <b>S</b> G	92.89	26CG	25031	26NE1	29.02
26CG	25031	67N	62.96	26CG	25031	163CB	84.79
26CG	25031	26CD2	9.01	26CG	25031	650	72.10
66CA	25031	65C	37.55	66CA	25031	65CA	60.47
66CA	25031	26CA	94.80	66CA	25031	26NE1	58.06
66CA	25031	67N	31.03	66CA	25031	26CD2	52.97
66CA	25031	650	32.59	65C	25031	65CA	22.96
65C	25031	26NE1	50.12	65C	25031	67N	68.57
65C	25031	26CD2	65.66	65C	25031	650	5.21
65CA	25031	26NE1	53.45	65CA	25031	67N	91.50
65CA	25031	26CD2	77.14	65CA	25031	650	28.05
26N	25031	26CA	20.41	2 <b>6N</b>	25031	25SG	49.78
26N	25031	26NE1	60.51	26N	25031	163CB	48.83
26N	25031	26CD2	54.87	26CA	25031	25SG	67.76
26CA	25031	26NE1	61.63	26CA	25031	67N	81.48
26CA		163CB	47.51	26CA	25031	26CD2	45.43
25SG	25031	26NE1	93.17	25 <b>SG</b>	25031		52.44
25NE1	25031	67N	75.15	26 <b>NE1</b>	25031	26CD2	27.61
26NE1	25031	650	51.59	67 <b>N</b>	25031	26CD2	55.94
67N	25031	650	63.57	163CB	25031	26CD2	92.09
26CD2	25031	650	64.59	25 <b>SG</b>	25 <b>N32</b>		82.28
25 <b>S</b> G	25N32	230	84.49	25 <b>S</b> G	25N32	26CD1	88.80
25 <i>S</i> G	25N32	26N	50.25	25 <b>SG</b>	25N32	25CB	9.66
25SG	25N32	25N	39.55	25 <b>S</b> G	25N32		46.38
25 <b>S</b> G	25N32	26CB	79.05	25SG	25N32	161C	82.48
25 <i>S</i> G	25N32	23C	74.25	1610	25N32	25CB	91.92

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			TAE	LE XVIII			
1610	25N32	163N	57.92	1610	25N32	161C	9.03
65CA	25N32	66N	32.94	65CA	25 <b>N3</b> 2	230	48.76
65CA	25N32	26CD1	52.31	65CA	25N32	26N	94.41
65CA	25N32	25N	92.40	65CA	25N32	65C	17.26
65CA	25N32	660	67.39	65CA	25N32	26CB	78.37
65CA	25N32	23C	60.93	66N	25N32	230	68.51
66N	25N32	26CD1	41.20	66N	25N32	26N	79.04
66N	25N32	25N	94.92	66N	25N32	65C	15.81
66N	25N32	660	35.14	6 <b>6</b> N	25N32	26CB	53.15
66N	25N32	23C	81.78	230	25N32	26CD1	46.25
230	25N32	26N	67.84	230	25N32	25CB	76.13
230	25N32	25N	47.96	230	25N32	65C	59.32
230	25N32	660	99.28	230	25N32	26CB	75.73
230	25N32	23C	13.32	26CD1	25N32	26N	42.44
26CD1	25N32	25CB	79.55	26CD1	25N32	25N	54.01
26CD1	25N32	65C	45.85	26CD1	25N32	660	<b>57.60</b>
26CD1	25N32	26CB	31.68	26CD1	25N32	23C	55.94
26N	25N32	25CB	42.72	26N	25N32	25 <b>N</b>	33.34
26N	25N32	65C	87.61	26N	25N32	660	74.76
26N	25N32	163N	67.13	26N	25N32	26CB	30.37
26N	25N32	23C	68.06	25CB	25N32	25N	30.07
25CB	25N32	163N	53.14	25CB	25N32	26CB	72.55
25CB	25N32	161C	92.10	25CB	25N32	23C	66.78
25N	25N32	65C	95.38	25N	25N32	163N	80.01
25N	25N32	26CB	60.55	25N	25N32	23C	41.68
65C	25N32	660	50.19	65C	25N32	26CB	65.71
65C	25N32	23C	72.49	660	25N32	163N	92.86
660	25N32	26CB	44.71	163N	25N32	26CB	78.55
163N	25N32	161C	52.17	26CB	25N32	23C	82.47
25 <i>S</i> G	25C33	25 <b>N</b>	62.03	25 <b>S</b> G	25C33	26N	67.44
25SG	25C33	25CB	22.53	25SG	25C33	25CA	44.14
25 <b>S</b> G	25C33	25C	51.85	25 <b>SG</b>	25C33	24N	94.55
25 <i>S</i> G	25C33	24C	75.69	25 <b>SG</b>	25C33	26CB	94.29
25 <b>S</b> G	25C33	24CA	93.12	25 <b>SG</b>	25C33	26CA	77.03
25SG	25C33	1610	66.31	230	25C33	25 <b>N</b>	68.18
230	25C33	26CD1	60.30	230	25C33	26N	92.61
230	25C33	23C	18.66	230	25C33	25CA	89.03
230	25C33	65CA	56.48	230	25C33	25C	96.38
230	25C33	24N	30.61	230	25C33	24C	56.80
230	25C33	26CG	75.69	230	25C33	26CB	94.54
230	25C33	23CA	32.16	230	25C33	66N	77.02
230	25C33	24CA	37.25	230	25C33	26CA	97.15
230	25C33	26NE1	47.95	230	25C33	65C	65.17
230	25C33	65N	43.20	25 <b>N</b>	25C33	26CD1	73.26

			TAB	LE XVIII			
25N	25C33	26N	44.51	25N	25C33	25CB	39.50
25N	25C33	23C	57.79	25N	25C33	25CA	20.88
25N	25C33	25C	34.96	25N	25C33	24N	40.75
25N	25C33	24C	13.79	25N	25C33	26CG	73.79
25N	25C33	26CB	75.59	25N	25C33	23CA	69.58
25N	25C33	24CA	31.77	25N	25C33	26CA	59.60
25N	25C33	26NE1	73.01	26CD1	25C33	26N	53.40
26CD1	25C33	23C	74.18	26CD1	25C33	25CA	82.03
26CD1	25C33	65CA	58.73	26CD1	25C33	25C	69.11
26CD1	25C33	24N	71.72	26CD1	25C33	24C	61.69
26CD1	25C33	26CG	15.97	26CD1	25C33	26CB	35.58
26CD1	25C33	23CA	91.94	26CD1	25C33	6 <b>6N</b>	43.00
26CD1	25C33	24CA	56.83	26CD1	25C33	26CA	46.45
26CD1	25C33	26NE1	12.72	26CD1	25C33	65C	47.83
26CD1	25C33	65N	61.48	26N	25C33	25CB	53.84
26N	25C33	23C	92.65	26N	25C33	25CA	37.01
26N	25C33	25C	17.36	26N	25C33	24N	77.69
26N	25C33	24C	45.23	26N	25C33	26CG	43.00
26N	25C33	26CB	34.40	26N	25C33	66N	88.08
26N	25C33	24CA	60.35	26N	25C33	26CA	15.70
26N	25C33	26NE1	62.43	26N	25C33	65C	98.83
25CB	25C33	23C	89.77	25CB	25C33	25CA	22.60
25CB	25C33	25C	36.53	25CB	25C33	24N	74.40
25CB	25C33	24C	53.16	25CB	25C33	26CG	96.54
25CB	25C33	26CB	86.56	25CB	25C33	23CA	92.73
25CB	25C33	24CA	70.80	25CB	25C33	26CA	67.26
25CB	25C33	1610	88.61	23C	25C33	25CA	78.15
23C	25C33	65CA	73.62	23C	25C33	25C	90.88
23C	25C33	24N	17.05	23C	25C33	24C	49.66
23C	25C33	26CG	88.11	23C	25C33	23CA	19.27
23C	25C33	66N	95.67	23C	25C33	24CA	32.32
23C	25C33	26NE1	62.90	23C	25C33	65C	83.62
23C	25C33	65N	59.43	25 <b>CA</b>	25C33	25C	21.13
25CA	25C33	24N	61.20	25CA	25C33	24C	33.01
25CA	25C33	26CG	76.92	25CA	25C33	26CB	71.41
25CA	25C33	23CA	87.72	25 <b>CA</b>	25C33	24CA	52.23
25CA	25033	26CA	52.44	25CA	25C33	26NE1	85.76
65CA	25C33	24N	87.00	65CA	25C33	26CG	69.63
65CA	25C33	26CB	84.46	65CA	25C33	23CA	76.25
65CA	25C33	66 <b>N</b>	32.22	65 <b>CA</b>	25C33	24CA	88.00
65CA	25C33	26NE1	51.85	65CA	25C33	65C	16.56
65CA	25C33	65N	14.77	2 =	25C33	24N	74.16
25C	25C33	2 <b>4C</b>	41.25	2:	25C33	26CG	60.10
25C	25C33	26CB	51.20	213	25C33	24CA	59.94

			TAB	LE XVIII			
25C	25C33	26CA	31.86	25C	25C33	26NE1	76.48
24N	25C33	24C	33.23	24N	25C33	26CG	82.78
24N	25C33	26CB	96.28	24N	25C33	23CA	32.00
24N	25C33	24CA	18.67	24N	25C33	26CA	88.81
24N	25C33	26NE1	62.87	24N	25C33	65C	93.98
24N	25C33	65N	73.77	24C	25C33	26CG	64.93
24C	25C33	26CB	70.95	24C	25C33	23CA	64.79
24C	25C33	24CA	19.55	24C	25C33	26CA	58.29
24C	25C33	26NE1	59.98	24C	25C33	65N	96.26
26CG	25C33	26CB	19.61	26CG	25C33	66N	46.12
26CG	25C33	24CA	65.68	26CG	25C33	26CA	32.66
26CG	25C33	26NE1	28.68	26CG	25C33	65C	55.83
26CG	25C33	65N	75.10	26CB	25C33	66N	55.47
26CB	25C33	24CA	77.83	26CB	25C33	26CA	19.41
26CB	25C33	26NE1	48.27	26CB	25C33	65C	68.78
26CB	25C33	65N	92.50	23CA	25C33	24CA	50.00
23CA	25C33	26NE1	79.91	23CA	25C33	65C	89.99
23CA	25C33	65N	61.53	6 <b>6 N</b>	25C33	24CA	94.59
66N	25C33	26CA	74.61	66N	25C33	26NE1	44.61
66N	25C33	65C	15.82	6 <b>6N</b>	25C33	65N	45.20
24CA	25C33	26CA	70.41	24CA	25C33	26NE1	50.59
24CA	25C33	65C	89.84	24CA	25C33	65N	77.59
26CA	25C33	26NE1	57.80	26CA	25C33	65C	87.10
26NE1	25C33	65C	44.55	26NE1	25C33	65N	51.65
65C	25C33	65N	29.48	660	25N34	66N	45.23
660	25N34	65CA	78.78	660	25N34	65C	62.40
660	25N34	66C	13.36	660	25N34	66CA	32.35
66N	25N34	65CA	34.26	66N	25N34	65C	17.63
66N	25N34	66C	35.00	6 <b>6N</b>	25N34	66CA	17.78
65CA	25N34	65C	20.48	65CA	25N34	66C	69.25
65CA	25N34	66CA	51.09	65C	25N34	66C	51.11
65C	25N34	66CA	31.81	66C	25N34	66CA	19.75

TABLE XIX

Table of angles between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one.

Atom 1	Atom 2	Atom 3	Angle	Atom 1	Atom 2	Atom 3	Angle
200	25C1	20C	8.55	200	25C1	21CA	35.54
200	25C1	20N	33.52	200	25C1	21N	19.74
200	25C1	21NE2	53.30	200	25C1	19CB	70.43
200	25C1	19CD	89.13	200	25C1	20CA	16.45
200	25C1	19CG	72.65	20C	25C1	21CA	32.39
20C	25C1	20N	30.42	20C	25C1	21N	14.60
20C	25C1	21NE2	58.84	20C	25C1	19CB	64.58
20C	25C1	19CD	81.24	20C	25C1	20CA	14.64
20C	25C1	19CG	65.09	184NE1	25C1	184CD1	18.88
184NE1	25C1	20N	99.03	184NE1	25C1	19CB	64.29
184NE1	25C1	19CD	56.93	184NE1	25C1	184CE2	13.44
184NE1	25C1	19CG	68.65	184CD1	25C1	20N	85.82
184CD1	25C1	19CB	57.54	184CD1	25C1	19CD	61.12
184CD1	25C1	184CE2	26.51	184CD1	25C1	19CG	67.61
21CA	25C1	20N	60.42	21CA	25C1	21N	18.12
21CA	25C1	21NE2	47.16	21CA	25C1	19CB	84.18
21CA	25C1	19CD	86.47	21CA	25C1	20CA	46.79
21CA	25C1	19CG	75.79	20N	25C1	21N	42.62
20N	25C1	21NE2	86.39	20 <b>N</b>	25C1	19CB	39.01
20N	25C1	19CD	64.17	20N	25C1	20CA	17.07
20N	25C1	19CG	47.30	21N	25C1	21NE2	54.26
21 <b>N</b>	25C1	19CB	71.03	21N	25C1	19CD	81.02
21N	25C1	20CA	28.72	21N	25C1	19CG	66.90
21NE2	25C1	20CA	69.50	19CB	25C1	19CD	29.32
19CB	25C1	20CA	54.83	19CB	25C1	184CE2	77.66
19CB	25C1	19CG	17.68	19CD	25C1	20CA	76.69
19CD	25C1	184CE2	69.09	19CD	25C1	19CG	17.08
20CA	25C1	19CG	<b>59.6</b> 5	184CE2	25C1	19CG	81.78
200	25C2	21NE2	59.99	200	25C2	20C	2.48
200	25C2	184NE1	99.98	200	25C2	184CD1	92.85

			TA	BLE XIX			
21NE2	25C2	2410H2	86.01	21NE2	25C2	20C	59.23
20C	25C2	184CD1	93.86	184NE1	25C2	184CD1	16.12
2410H2	25C3	21NE2	97.71	200	25C3	21NE2	54.27
200	25C3	184CD1	87.74	200	25C3	20C	3.17
21NE2	25C3	20C	54.18	184CD1	25C3	20C	88.87
2410H2	25C4	1840	97.32	200	25C4	180D1	66.74
200	25C4	184CD1	99.13	200	25C4	18CG	78.46
200	25C4	18ND2	91.26	200	25C4	20C	7.41
200	25C4	184NE1	94.68	200	25C4	20N	34.24
180D1	25C4	184CD1	86.06	180D1	25C4	184CG	91.60
180D1	25C4	1840	62.83	180D1	25C4	184CB	82.77
180D1	25C4	184CA	64.10	180D1	25C4	18CG	12.46
180D1	25C4	18ND2	28.21	180D1	25C4	20C	59.47
180D1	25C4	184NE1	97.79	180D1	25C4	20 <b>N</b>	35.84
180D1	25C4	184C	57.15	184CD1	25C4	184CG	17.48
184CD1	25C4	1840	65.93	184CD1	25C4	184CB	33.13
184CD1	25C4	184CA	37.60	184CD1	25C4	18CG	88.07
184CD1	25C4	18ND2	95.74	184CD1	25C4	20C	99.37
184CD1	25C4	184NE1	15.35	184CD1	25C4	20N	82.14
184CD1	25C4	184C	55.11	184CG	25C4	1840	54.64
184CG	25C4	184CB	18.72	184CG	25C4	184CA	31.21
184CG	25C4	18CG	89.91	184CG	25C4	18ND2	92.98
184CG	25C4	184NE1	27.12	184CG	25C4	20N	96.33
184CG	25C4	184C	46.92	1840	25C4	184CB	36.11
1840	25C4	184CA	29.52	1840	25C4	18CG	52.77
1840	25C4	18ND2	45.59	1840	25C4	184NE1	80.28
1840	25C4	20N	93.80	1840	25C4	184C	13.81
184CB	25C4	184CA	18.71	184CB	25C4	18CG	77.83
184CB	25C4	18ND2	77.39	184CB	25C4	184NE1	45.48
184CB	25C4	20N	98.33	184CB	25C4	184C	30.19
184CA	25C4	18CG	59.86	184CA	25C4	18ND2	61.77
184CA	25C4	184NE1	52.74	184CA	25C4	20N	81.89
184CA	25C4	184C	17.52	18CG	25C4	18ND2	16.14
18CG	25C4	20C	71.10	18CG	25C4	20 <b>N</b>	48.30
18CG	25C4	184C	49.48	18ND2	25C4	20C	83.86
18ND2	25C4	20N	63.67	18ND2	25C4	184C	47.21
20C	25C4	184NE1	96.84	200	25C4	20N	28.14
184NE1	25C4	20N	86.08	184NE1	25C4	184C	70.17

			TA	BLE XIX			
20N	25C4	184C	83.78	200	25C5	180D1	80.82
200	25C5	20N	45.03	200	25C5	20C	11.42
200	25C5	18CG	91.75	200	25C5	19CB	74.81
200	25C5	20CA	30.47	200	25C5	19N	82.23
200	25C5	19C	51.72	200	25C5	19CA	69.30
180D1	25C5	20N	46.49	180D1	25C5	184CA	77.31
180D1	25C5	20C	70.82	180D1	25C5	18CG	11.80
180D1	25C5	184CB	96.95	180D1	25C5	19CB	72.36
180D1	25C5	1830	66.28	180D1	25C5	1840	67.87
180D1	25C5	20CA	51.82	180D1	25C5	19N	42.94
180D1	25C5	184C	64.99	180D1	25C5	18ND2	27.18
180D1	25C5	19C	53.47	180D1	25C5	19CA	55.00
184CD1	25C5	184CG	18.87	184CD1	25C5	184NE1	17.79
184CD1	25C5	184CA	44.65	184CD1	25C5	184CB	36.67
184CD1	25C5	19CB	67.07	184CD1	25C5	1830	47.98
184CD1	25C5	1840	72.91	184CD1	25C5	19 <b>N</b>	80.31
184CD1	25C5	184C	63.25	184CD1	25C5	19C	98.13
184CD1	25C5	19CA	81.55	184CD1	25C5	184CD2	22.19
20N	25C5	20C	33.70	20N	25C5	18CG	58.10
20N	25C5	19CB	44.66	20N	25C5	1830	72.78
20N	25C5	20CA	18.20	20N	25C5	19N	38.00
20N	25C5	18ND2	73.36	20N	25C5	19C	13.46
20N	25C5	19CA	30.03	184CG	25C5	184NE1	31.64
184CG	25C5	184CA	35.78	184CG	25C5	184CB	20.39
184CG	25C5	19CB	83.44	184CG	25C5	1830	54.59
184CG	25C5	1840	58.05	184CG	25C5	19N	89.86
184CG	25C5	184C	51.94	184CG	25C5	19CA	95.52
184CG	25C5	184CD2	13.48	184NE1	25C5	184CA	62.33
184NE1	25C5	184CB	51.80	184NE1	25C5	19CB	68.57
184NE1	25C5	1830	61.84	184NE1	25C5	1840	89.24
184NE1	25C5	19N	89.54	184NE1	25C5	184C	80.74
184NE1	25C5	19C	<b>39.</b> 19	184NE1	25C5	19CA	85.71
184NE1	25C5	184CD2	26.19	184CA	25C5	18CG	69.03
184CA	25C5	184CB	20.92	18 <b>4CA</b>	25 <b>C</b> 5	19CB	77.96
184CA	25C5	1830	36.00	18 <b>4CA</b>	25C5	1840	31.59
184CA	25C5	19N	56.93	184CA	25C5	184C	18.88
184CA	25C5	18ND2	67.71	184CA	25C5	19C	97.51
184CA	25C5	19CA	80.33	184CA	25C5	184CD2	49.00

			TA	BLE XIX			
20C	25C5	18CG	82.15	20C	25C5	19CB	67.05
20C	25C5	20CA	19.43	20C	25C5	19N	71.23
20C	25C5	18ND2	92.73	20C	25C5	19C	41.28
20C	25C5	19CA	59.31	18CG	25C5	184CB	87.50
18CG	25C5	19CB	79.43	18CG	25C5	1830	64.87
18CG	25C5	1840	56.38	18CG	25C5	20CA	63.40
18CG	25C5	19N	48.70	18CG	25C5	184C	54.85
18CG	25C5	18ND2	16.71	18CG	25C5	19C	64.20
18CG	25C5	19CA	63.20	184CB	25C5	19CB	90.65
184CB	25C5	1830	52.34	184CB	25C5	1840	37.66
184CB	25C5	19N	86.29	184CB	25C5	184C	32.87
184CB	25C5	18ND2	82.52	184CB	25C5	19CA	97.44
184CB	25C5	2410H2	93.49	184CB	25C5	184CD2	31.41
19CB	25C5	1830	42.37	19CB	25C5	20CA	60.88
19CB	25C5	19N	30.92	19CB	25C5	184C	88.12
19CB	25C5	18ND2	95.45	19CB	25C5	19C	31.26
19CB	25C5	19CA	18.10	19CB	25C5	184CD2	89.12
1830	25C5	1840	61.91	1830	25C5	20CA	90.84
1830	25C5	19N	35.42	1830	25C5	184C	46.49
1830	25C5	18ND2	74.22	1830	25C5	19C	63.18
1830	25C5	19CA	45.18	1830	25C5	184CD2	66.30
1840	25C5	19N	81.26	1840	25C5	184C	15.43
1840	25C5	18ND2	46.67	1840	25C5	19CA	98.50
1840	25C5	2410H2	80.65	1840	25C5	184CD2	68.07
20CA	25C5	19N	55.69	20CA	25C5	18ND2	75.55
20CA	25C5	19C	30.17	20CA	25C5	19CA	47.94
19N	25C5	184C	68.55	19N	25C5	18ND2	64.53
19N	25C5	19C	31.12	19N	25C5	19CA	17.72
184C	25C5	18ND2	50.25	184C	25C5	19C	99.32
184C	25C5	19CA	85.02	184C	25C5	2410H2	95.85
184C	25C5	184CD2	64.13	18ND2	25C5	19C	80.57
18ND2	25C5	19CA	79.75	18ND2	25C5	2410H2	86.00
19C	25C5	19CA	18.28	241CH2	25 <b>C</b> 5	184CD2	93.94
200	25C6	20C	12.91	200	25C6	20N	48.34
200	25C6	19CB	91.47	200	25C6	180D1	72.52
200	25C6	19CG	86.64	200	25 <b>C</b> 6	20CA	29.66
200	25C6	19C	58.59	200	25 <b>C6</b>	19CA	78.30
200	25C6	19N	85.64	200	25C6	21N	15.81

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			TA	BLE XIX			
200	25C6	19NE2	96.10	184CD1	25C6	184NE1	21.76
184CD1	25C6	19CB	75.50	184CD1	25C6	180D1	95.54
184CD1	25C6	19CG	83.68	184CD1	25C6	184CG	14.32
184CD1	25C6	19CD	72.31	184CD1	25C6	19CA	87.17
184CD1	25C6	190E1	56.50	184CD1	25C6	19N	79.17
184CD1	25C6	184CE2	25.40	184CD1	25C6	1830	45.26
184CD1	25C6	184CA	35.59	184CD1	25C6	184CB	27.68
184CD1	25C6	19NE2	80.74	184NE1	25C6	19CB	80.25
184NE1	25C6	19CG	80.79	184NE1	25C6	184CG	30.82
184NE1	25C6	19CD	64.39	184NE1	25C6	19CA	96.31
184NE1	25C6	190E1	49.89	184NE1	25C6	19N	93.83
184NE1	25C6	184CE2	11.71	184NE1	25C6	1830	62.24
184NE1	25C6	184CA	57.31	184NE1	25C6	184CB	47.06
184NE1	25C6	19NE2	68.19	20C	25C6	20N	37.04
20C	25C6	19CB	78.57	20C	25C6	180D1	67.96
20C	25C6	19CG	74.74	20C	25C6	20CA	19.73
20C	25C6	19C	45.89	20C	25C6	19CD	90.64
20C	25C6	19CA	65.64	20C	25C6	19N	74.27
20C	25C6	21N	11.50	20C	25C6	19NE2	87.59
20N	25C6	19CB	50.10	20N	25C6	180D1	42.87
20N	25C6	19CG	57.53	20N	25C6	20CA	19.02
20N	25C6	19C	16.56	20N	25C6	19CD	76.93
20N	25C6	19CA	32.67	20N	25C6	190E1	86.01
20N	25C6	19N	37.35	20N	25C6	1830	71.42
20N	25C6	21N	45.38	20N	25C6	184CA	93.41
20N	25C6	19NE2	82.82	19CB	25C6	180D1	71.17
19CB	25C6	19CG	20.52	19CB	25C6	184CG	87.09
19CB	25C6	20CA	67.16	19CB	25C6	19C	34.76
19CB	25C6	19CD	33.99	19CB	25C6	19CA	18.63
19CB	25C6	190E1	37.17	19CB	25C6	19N	30.79
19CB	25C6	184CE2	91.78	19CB	25C6	1830	42.56
19CB	25C6	21N	80.11	19CB	25C6	184CA	74.81
19CB	25C6	184CB	88.09	19CB	25 <b>C</b> 6	19NE2	46.70
180D1	25C6	19CG	88.69	180D1	25C6	184CG	91.56
180D1	25C6	20 <b>CA</b>	49.91	180 <b>D1</b>	25C6	19C	53.85
180D1	25C6	19CA	54.76	180 <b>D1</b> .	25 <b>C</b> 6	19N	40.79
180D1	25C6	1830	58.55	180 <b>D1</b>	25 <b>C6</b>	21N	79.33
180D1	25C6	184CA	61.28	180D1	25 <b>C</b> 6	184CB	76.32

			TA	BLE XIX			
19CG	25C6	184CG	97.34	19CG	25C6	20CA	70.01
19CG	25C6	19C	41.04	19CG	25C6	19CD	19.41
19CG	25C6	19CA	33.94	19CG	25C6	190E1	31.04
19CG	25C6	19N	50.11	19CG	25C6	184CE2	92.31
19CG	25C6	1830	60.89	19CG	25C6	21N	72.24
19CG	25C6	184CA	92.44	19CG	25C6	19NE2	28.22
184CG	25C6	19CD	86.63	184CG	25C6	19CA	96.09
184CG	25C6	190E1	70.82	184CG	25C6	19N	84.76
184CG	25C6	184CE2	28.58	184CG	25C6	1830	51.08
184CG	25C6	184CA	30.50	184CG	25C6	184CB	16.46
184CG	25C6	19NE2	94.82	20CA	25C6	19C	32.41
20CA	25C6	19CD	88.73	20CA	25C6	19CA	50.99
20CA	25C6	19N	56.06	20CA	25C6	1830	89.93
20CA	25C6	21N	30.16	20CA	25 <b>C</b> 6	19NE2	90.65
19C	25C6	19CD	60.43	19C	25C6	19CA	19.75
19C	25C6	190E1	69.63	19C	25C6	19N	32.09
19C	25C6	1830	64.40	19C	25 <b>C</b> 6	21N	50.78
19C	25C6	184CA	92.13	19C	25C6	19NE2	67.14
19CD	25C6	19CA	51.19	19CD	25C6	190E1	15.82
19CD	25C6	19N	64.65	19CD	25C6	184CE2	75.35
19CD	25C6	1830	64.06	19CD	25C6	21N	85.72
19CD	25C6	184CA	91.40	19CD	25C6	184CB	96.57
19CD	25C6	19NE2	14.56	19CA	25C6	190E1	55.80
19CA	25C6	19 <b>N</b>	18.35	19CA	25C6	1830	45.71
19CA	25C6	21N	70.13	19CA	25C6	184CA	75.80
19CA	25C6	184CB	92.14	19CA	25C6	19NE2	62.13
190E1	25C6	19N	64.68	190E1	25C6	184CE2	61.31
190E1	25C6	1830	54.29	190E1	25C6	184CA	77.83
190E1	25C6	184CB	81.20	190E1	25C6	19NE2	26.71
19N	25C6	1830	34.10	19N	25C6	21N	81.52
19N	25C6	184CA	60.05	19N	25C6	184CB	77.37
19N	25C6	19NE2	77.44	184CE2	25C6	1830	69.91
184CE2	25C6	184CA	58.41	184CE2	25C6	184CB	44.79
184CE2	25C6	19NE2	77.70	1830	25C6	184CA	32.25
1830	25C6	184CB	46.82	1830	25C6	19NE2	78.49
21N	25C6	19NE2	80.31	184CA	25C6	184CB	17.72
200	2507	21.CA	42.87	200	2507	20C	13.38
200	2507	21C	58.81	200	2507	19CD	93.04

	TABLE XIX								
200	2507	210	73.65	200	2507	220	77.10		
200	2507	21N	27.93	200	2507	19CG	74.84		
200	2507	190E1	98.59	200	2507	184CD1	93.22		
200	2507	19CB	66.24	21CA	2507	20C	34.50		
21CA	2507	19NE2	96.98	21CA	2507	21C	20.45		
21CA	2507	19CD	99.50	21CA	2507	210	32.06		
21CA	2507	220	59.78	21CA	2507	21N	18.66		
21CA	2507	19CG	84.02	21CA	2507	19CB	87.69		
20C	2507	19NE2	92.76	20C	2507	21C	47.32		
20C	2507	19CD	85.47	20C	2507	210	62.79		
20C	2507	220	64.15	20C	2507	21N	16.77		
20C	2507	19CG	67.03	20C	2507	190E1	93.62		
20C	2507	184CD1	98.91	20C	2507	19CB	62.11		
19NE2	2507	21C	79.95	19NE2	2507	19CD	17.34		
19NE2	2507	184NE1	69.42	19NE2	2507	210	80.15		
19NE2	2507	220	37.30	19NE2	2507	21N	90.94		
19NE2	2507	19CG	30.25	19 <b>NE</b> 2	2507	190E1	27.55		
19NE2	2507	184CD1	74.17	19 <b>NE</b> 2	2507	19CB	45.70		
21C	2507	19CD	86.46	21C	2507	210	15.78		
21C	2507	220	43.07	21C	2507	21N	30.95		
21C	2507	19CG	74.61	21C	2507	19CB	83.86		
19CD	2507	184NE1	58.81	19CD	2507	210	90.75		
19CD	2507	220	44.18	19CD	2507	21N	88.47		
19CD	2507	19CG	18.45	19CD	2507	190E1	14.37		
19CD	2507	184CD1	60.00	19CD	2507	19CB	29.83		
184NE1	2507	19CG	<b>69</b> .06	184NE1	2507	190E1	44.47		
184NE1	2507	184CD1	15.81	184NE1	2507	19CB	60.87		
210	2507	220	46.84	210	2507	21N	46.16		
210	2507	19CG	82.83	210	2507	19CB	95.13		
220	2507	21N	56.64	220	2507	19CG	37.32		
220	2507	190E1	58.55	220	2507	19CB	52.98		
21N	2507	19CG	70.99	21N	2507	190E1	99.59		
21N	2507	19CB	71.16	19CG	2507	190E1	28.77		
19CG	2507	184CD1	65.14	19CG	2507	19CB	17.77		
190E1	2507	184CD1	46.81	190E1	2507	19CB	32.36		
184CD1	2507	19CB	52.79	19NE2	25 <b>C8</b>	184NE1	74.20		
19NE2	25C8	19CD	17.18	19NE2	25C8	200	86.21		
19NE2	25C8	190E1	28.35	19NE2	25C8	220	35.57		

			TA	BLE XIX			
19NE2	25C8	210	77.21	19NE2	25C8	184CE2	85.22
184NE1	25C8	19CD	60.55	184NE1	25C8	200	90.70
184NE1	25C8	190E1	46.23	184NE1	25C8	184CE2	14.61
19CD	25C8	200	76.90	19CD	25C8	190E1	15.08
19CD	25C8	220	42.30	19CD	25C8	210	84.98
19CD	25C8	184CE2	73.32	200	25C8	190E1	83.60
200	25C8	220	62.93	200	25C8	210	59.11
190E1	25C8	220	57.39	190E1	25C8	184CE2	58.44
220	25C8	210	42.86	184NE1	25C9	19CD	81.47
184NE1	25C9	190E1	63.24	184NE1	25C9	184CE2	17.78
184NE1	25C9	184CD1	15.24	184NE1	25C9	184CZ2	36.54
184NE1	25C9	19CG	84.02	184NE1	25C9	162NE2	48.91
184NE1	25 <b>C</b> 9	162CD2	64.08	184NE1	25C9	19CB	68.93
19NE2	25C9	19CD	22.29	19NE2	25C9	190E1	38.48
19NE2	25C9	184CD1	96.98	19NE2	25C9	19CG	31.33
19NE2	25C9	162NE2	73.61	19NE2	25C9	220	33.83
19NE2	25C9	162CD2	72.04	19NE2	25C9	19CB	48.00
19CD	25C9	190E1	20.30	19CD	25C9	184CE2	96.87
19CD	25C9	184CD1	75.05	19CD	25C9	19CG	16.69
19CD	25C9	162NE2	65.70	19CD	25C9	220	43.91
19CD	25C9	162CD2	70.64	19CD	25C9	19CB	28.29
190E1	25C9	184CE2	77.36	190E1	25 <b>C9</b>	184CD1	59.86
190E1	25C9	184CZ2	88.34	190E1	25 <b>C</b> 9	19CG	32.36
190E1	25C9	162NE2	47.50	190E1	25C9	220	64.20
190E1	25C9	162CD2	55.96	190E1	25C9	19CB	33.06
184CE2	25C9	184CD1	31.58	184CE2	25C9	184CZ2	19.41
184CE2	25C9	162NE2	49.67	184CE2	25 <b>C9</b>	162CD2	61.51
184CE2	25C9	19CB	86.67	184CD1	25C9	184CZ2	50.94
184CD1	25C9	19CG	73.85	184CD1	25C9	162NE2	59.35
184CD1	25C9	162CD2	75.48	184CD1	25C9	19CB	57.22
184CZ2	25 <b>C</b> 9	162NE2	48.81	184CZ2	25 <b>C9</b>	162CD2	55.09
19CG	25C9	162NE2	79.83	19CG	25C9	220	36.66
19CG	25C9	162CD2	86.65	19CG	25 <b>C9</b>	19CB	17.77
162NE2	25C9	162CD2	16.38	162NE2	25 <b>C9</b>	19CB	77.35
220	25C9	19CB	51.74	162CD2	25 <b>C9</b>	19CB	88.41
184NE1	25C10	184CZ2	41.93	184NE1	25C10	19NE2	89.55
184NE1	25C10	184CE2	20.67	1.84NE1	25C10	162CD2	76.32
184NE1	25C10	190E1	57.76	184NE1	25C10	162NE2	56.54

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		TA	BLE XIX			
184NE1	25C10 19CD	70.87	184NE1	25C10	184CD1	8.43
184NE1	25C10 162CG	77.88	184NE1	25C10	184CH2	49.27
184NE1	25C10 162CE1	51.80	184CZ2	25C10	184CE2	21.41
184CZ2	25C10 162CD2	68.85	184CZ2	25C10	190E1	92.45
184CZ2	25C10 162NE2	58.15	184CZ2	25C10	184CD1	50.03
184CZ2	25C10 162CG	61.13	184CZ2	25C10	184CH2	7.67
184CZ2	25C10 162CE1	46.19	19NE2	25C10	162CD2	81.68
19NE2	25C10 19OE1	34.40	19NE2	25C10	162NE2	79.07
19NE2	25C10 19CD	18.69	19NE2	25C10	184CD1	83.31
19NE2	25C10 162CG	94.78	19NE2	25C10	162CE1	90.70
184CE2	25C10 162CD2	73.19	184CE2	25C10	190E1	75.46
184CE2	25C10 162NE2	56.50	184CE2	25C10	19CD	90.32
184CE2	25C10 184CD1	28.64	184CE2	25C10	162CG	70.03
184CE2	25C10 184CH2	28.62	184CE2	25C10	162CE1	47.27
162CD2	25C10 19OE1	63.96	162CD2	25C10	162NE2	20.18
162CD2	25C10 19CD	77.09	162CD2	25C10	184CD1	81.08
162CD2	25C10 162CG	13.20	162CD2	25C10	184CH2	71.00
162CD2	25C10 162CE1	25.99	190E1	25C10	162NE2	52.28
190E1	25C10 19CD	17.69	190E1	25C10	184CD1	53.34
190E1	25C10 162CG	76.09	190E1	25C10	184CH2	99.88
190E1	25C10 162CE1	61.59	162NE2	25C10	19CD	68.59
162NE2	25C10 184CD1	60.95	162NE2	25C10	162CG	26.84
162NE2	25C10 184CH2	62.65	162NE2	25C10	162CE1	12.34
19CD	25C10 184CD1	64.80	19CD	25C10	162CG	90.03
19CD	25C10 162CE1	78.73	184CD1	25C10	162CG	84.10
184CD1	25C10 184CH2	57.23	184CD1	25C10	162CE1	57.81
162CG	25C10 184CH2	61.82	162CG	25C10	162CE1	26.29
184CH2	25C10 162CE1	50.38	18 <b>4CZ2</b>	25C11	184NE1	33.44
184CZ2	25C11 162CD2	58.33	18 <b>4CZ2</b>	25C11	184CE2	16.76
184CZ2	25C11 19NE2	94.98	184CZ2	25C11	162NE2	47.17
184NE1	25C11 162CD2	59.46	184NE1	25C11	184CE2	17.00
184NE1	25C11 19NE2	<b>55.</b> 09	184NE1	25C11	162NE2	43.40
162CD2	25C11 184CE2	5 <b>9</b> .69	162CD2	25C11	19NE2	64.20
162CD2	25C11 162NE2	16.09	184CE2	25C11	19NE2	81.10
184CE2	25C11 162NE2	45.09	19NE2	25011	162NE2	60.40
162CD2	25S14 184CZ2	63.39	162CD2	25514	162CG	18.68
162CD2	25S14 162CB	3€.19	162CD2	25514	1610	65.32
162CD2	25S14 162NE2	15.75	162CD2	25514	1610D1	86.18

			TAI	BLE XIX			
162CD2	25514	184CH2	70.43	184CZ2	25\$14	162CG	63.65
184CZ2	<b>25S14</b>	162CB	78.78	184CZ2	25514	162NE2	49.18
184CZ2	25514	184CH2	14.76	162CG	25S14	162CB	20.04
162CG	<b>25S14</b>	1610	60.62	162CG	25S14	162NE2	28.04
162CG	25S14	1610D1	69.44	162CG	25\$14	184CH2	65.75
162CB	25514	1610	46.23	162CB	25\$14	162NE2	47.97
162CB	25514	1610D1	50.06	162CB	25514	184CH2	76.80
1610	25514	162NE2	81.07	1610	25514	1610D1	48.28
162NE2	25S14	1610D1	97.37	162NE2	25S14	184CH2	58.34
1610D1	25\$14	184CH2	95.88	184CZ2	25015	184CH2	22.20
184CZ2	25015	162CG	77.78	184CZ2	25015	162CD2	73.98
184CZ2	25015	1370	88.43	184CZ2	25015	162CB	96.58
184CZ2	25015	184CE2	11.44	184CZ2	25015	137CB	79.42
184CZ2	25015	162ND1	62.04	184CZ2	25015	162NE2	56.17
184CZ2	25015	162CE1	49.68	184CZ2	25015	137C	97.15
184CZ2	25015	184NE1	27.38	184CZ2	25015	184CZ3	22.14
184CH2	25015	162CG	83.63	184CH2	25015	162CD2	86.91
184CH2	25015	1370	66.28	184CH2	25015	162CB	97.10
184CH2	25015	184CE2	33.64	184CH2	25015	137CB	63.34
184CH2	25015	162ND1	65.67	184CH2	25015	162NE2	70.78
184CH2	25015	162CE1	59.07	184CH2	25015	137C	75.65
184CH2	25015	184NE1	49.58	184CH2	25015	184CZ3	0.44
162CG	25015	162CD2	20.14	162CG	25015	1370	98.89
162CG	25015	162CB	21.58	162CG	25015	184CE2	75.94
162CG	25015	137CB	60.62	162CG	25015	162ND1	18.08
162CG	25015	162NE2	30.14	162CG	25015	1610D1	74.08
162CG	25015	162CE1	28.30	162CG	25015	137C	90.40
162CG	25015	184NE1	73.88	162CG	25015	184CZ3	84.03
162CD2	25015	162CB	38.15	162CD2	25015	184CE2	68.37
162CD2	25015	137CB	79.85	162CD2	25015	162ND1	30.08
162CD2	25015	162NE2	17.97	162CD2	25015	1610D1	89.07
162CD2	25015	162CE1	28.23	162CD2	25015	184NE1	61.40
162CD2	25015	184CZ3	87.22	137C	25015	162CB	88.99
1370	25015	184CE2	99.87	1370	25015	137CB	38.31
1370	25015	162ND1	89.22	1370	25015	1610D1	64.44
1370	25015	162CE1	99.08	1370	25015	137C	13.71
1370	25015	184CZ3	66.38	162CB	25015	184CE2	96.47
162CB	25015	137CB	53.88	162CB	25015	162ND1	34.56

		TA	BLE XIX		·
162CB	25015 162NE2	51.49	162CB	25015 1610D1	52.55
162CB	25015 162CE1	48.77	162CB	25015 137C	77.54
162CB	25015 184NE1	95.45	162CB	25015 184CZ3	97.53
184CE2	25015 137CB	88.43	184CE2	25015 162ND1	62.44
184CE2	25015 162NE2	50.44	184CE2	25015 162CE1	47.79
184CE2	25015 184NE1	15.94	184CE2	25015 184CZ3	33.57
137CB	25015 162ND1	52.03	137CB	25015 162NE2	80.03
137CB	25015 1610D1	58.03	137CB	25015 162CE1	64.80
137CB	25015 137C	32.11	137CB	25015 184CZ3	63.70
162ND1	25015 162NE2	28.43	162ND1	25015 1610D1	83.20
162ND1	25015 162CE1	16.55	162ND1	25015 137C	84.06
162ND1	25015 184NE1	64.50	162ND1	25015 184CZ3	66.08
162NE2	25015 162CE1	16.59	162NE2	25015 184NE1	44.67
162NE2	25015 184CZ3	71.04	1610D1	25015 162CE1	99.43
1610D1	25015 137C	51.29	162CE1	25015 137C	96.68
162CE1	25015 184NE1	48.13	162CE1	25015 184CZ3	59.41
137C	25015 184CZ3	75.83	184NE1	25015 184CZ3	49.51
1610D1	25016 1610	49.89	1610D1	25016 161CG	14.03
1610	25016 161CG	47.85	162CD2	25N17 162CG	24.82
162CD2	25N17 1610	98.82	162CD2	25N17 162CB	49.71
162CD2	25N17 162CA	57.32	162CD2	25N17 162NE2	15.00
162CD2	25N17 161C	92.83	162CD2	25N17 25SG	58.11
162CD2	25N17 162N	75.97	162CD2	25N17 162ND1	25.66
162CD2	25N17 162CE1	19.01	162CD2	25N17 184CZ2	61.54
162CD2	25N17 25CB	41.82	162CG	25N17 1610	86.86
162CG	25N17 162CB	26.56	162CG	25N17 162CA	42.37
162CG	25N17 162NE2	34.27	162CG	25N17 161C	76.28
162CG	25N17 25SG	73.91	162CG	25N17 162N	58.18
162CG	25N17 162ND1	11.86	162CG	25N17 1610D1	80.11
162CG	25N17 162CE1	24.71	162CG	25N17 184CZ2	63.21
162CG	25N17 25CB	63.12	1610	25N17 162CB	64.87
1610	25N17 162CA	44.51	1610	25N17 161C	15.53
1610	25N17 25SG	72.13	1610	25N17 162N	31.00
1610	25N17 162ND1	97.99	1610	25N17 1610D1	55.13
1610	25N17 25CB	90.74	162CB	25N17 162CA	24.47
162CB	25N17 162NE2	60.78	162CB	25N17 161C	51.89
152CB	25N17 25SG	82.14	162CB	25N17 162N	34.09
162CB	25N17 162ND1	34.92	162CB	25N17 1610D1	55.89

			TA	BLE XIX			
162CB	25N17	162CE1	50.58	162CB	25N17	184CZ2	81.16
162CB	25N17	25CB	79.93	162CA	25N17	162NE2	71.74
162CA	25N17	161C	35.58	162CA	25N17	25SG	65.34
162CA	25N17	162N	19.04	162CA	25N17	162ND1	53.71
162CA	25N17	1610D1	60.49	162CA	25N17	162CE1	66.39
162CA	25N17	25CB	71.10	162NE2	25N17	25SG	65.94
162NE2	25N17	162N	89.98	162NE2	25N17	162ND1	29.38
162NE2	25N17	162CE1	14.46	162NE2	25N17	184CZ2	48.97
162NE2	25N17	25CB	46.17	161C	25N17	25SG	80.41
161C	25N17	162N	18.12	161C	25N17	162ND1	86.43
161C	25N17	1610D1	42.82	161C	25N17	25CB	95.81
25SG	25N17	162N	77.28	25SG	25N17	162ND1	81.39
25SG	25 <b>N17</b>	162CE1	76.71	25SG	25N17	25CB	21.56
162N	25N17	162ND1	68.34	162N	25N17	1610D1	45.06
162N	25N17	162CE1	82.88	162N	25N17	25CB	87.52
162ND1	25N17	1610D1	83.28	162ND1	25N17	162CE1	16.35
162ND1	25 <b>N17</b>	184CZ2	51.35	162ND1	25N17	25CB	67.28
1610D1	25N17	162CE1	99.02	1610D1	25N17	184CZ2	99.72
162CE1	25 <b>N</b> 17	184CZ2	43.80	162CE1	25N17	25CB	58.69
184CZ2	25N17	25CB	91.04	1610	25C18	25 <b>S</b> G	88.17
1610	25C18	162CD2	89.98	1610	25C18	162CA	40.67
1610	25C18	161C	9.73	1610	25C18	162CG	74.76
1610	25C18	162CB	55.30	1610	25C18	162N	24.42
25SG	25C18	162CD2	63.21	25 <b>S</b> G	25C18	162CA	69.19
25SG	25C18	161C	89.73	25SG	25C18	162CG	72.88
25 <i>S</i> G	25C18	162CB	81.01	25 <b>SG</b>	25C18	25CB	22.17
25 <i>S</i> G	25C18	19NE2	60.90	25 <b>SG</b>	25C18	162NE2	64.84
25SG	25C18	162N	81.28	162CD2	25C18	162CA	49.41
162CD2	25C18	161C	82.11	162CD2	25C18	162CG	17.90
162CD2	25C18	162CB	38.19	162CD2	25C18	25CB	48.19
162CD2	25C18	19NE2	74.06	1.62CD2	25C18	162NE2	11.85
162CD2	25C18	162N	65.91	162CA	25C18	161C	33.66
162CA	25C18	162CG	35.77	162CA	25C18	162CB	21.19
162CA	25C18	25CB	73.11	162CA	25C18	162NE2	61.26
162CA	25C18	162N	17.67	161C	25C18	162CG	66.10
161C	25C18	162CB	46.17	161C	25C18	162NE2	93.83
161C	25C18	162N	16.25	162CG	25C18	162CB	20.62
162CG	25C18	25CB	62.56	162CG	25C18	19NE2	91.91

			TA	BLE XIX			
162CG	25C18	162NE2	28.46	162CG	25C18	162N	50.34
162CB	25C18	25CB	77.17	162CB	25C18	162NE2	49.07
162CB	25C18	162N	31.21	25CB	25C18	19NE2	46.45
25CB	25C18	162NE2	46.22	25CB	25C18	162N	89.13
19NE2	25C18	162NE2	63.90	162NE2	25C18	162N	77.68
25 <b>S</b> G	25C19	25CB	31.14	25SG	25C19	162CD2	72.89
25SG	25C19	1610	97.36	25 <b>S</b> G	25C19	19NE2	87.55
25SG	25C19	25N	42.09	25 <i>S</i> G	25C19	25CA	29.35
25SG	25C19	23CA	99.22	25 <b>S</b> G	25C19	162CA	71.95
25SG	25C19	230	76.04	25 <b>S</b> G	25C19	23C	80.42
25 <b>S</b> G	25C19	162CG	75.66	25 <b>S</b> G	25C19	162NE2	71.54
25SG	25C19	19CD	78.54	25 <b>S</b> G	25C19	190E1	73.25
25SG	25C19	161C	92.06	25 <i>S</i> G	25C19	162CB	81.06
25CB	25C19	162CD2	57.89	25CB	25C19	19NE2	60.82
25CB	25C19	25N	33.86	25CB	25C19	25CA	15.34
25CB	25C19	23CA	90.78	25CB	25C19	162CA	81.46
25CB	25C19	230	81.85	25CB	25C19	23C	77.61
25CB	25C19	162CG	67.35	25CB	25C19	162NE2	50.15
25CB	25C19	19CD	49.61	25CB	25C19	190E1	42.26
25CB	25C19	162CB	81.00	162CD2	25C19	1610	78.28
162CD2	25C19	19NE2	84.27	162CD2	25C19	25N	90.80
162CD2	25C19	25CA	73.23	162CD2	25C19	162CA	45.80
162CD2	25C19	162CG	13.99	162CD2	25C19	162NE2	13.84
162CD2	25C19	19CD	72.50	162CD2	25C19	190E1	57.48
162CD2	25C19	161C	71.45	162CD2	25C19	162CB	32.02
1610	25C19	162CA	36.90	1610	25C19	1.62CG	64.31
1610	25C19	162NE2	92.09	1610	25C19	161C	7.56
1610	25C19	162CB	46.27	19NE2	25C19	25N	50.87
19NE2	25C19	25CA	58.20	19NE2	25C19	23CA	48.05
19NE2	25C19	230	68.25	19NE2	25C19	23C	52.78
19NE2	25C19	162CG	98.10	19NE2	25C19	162NE2	70.55
19NE2	25C19	19CD	13.21	19NE2	25C19	190El	27.94
25N	25C19	25CA	19.24	25N	25C19	23CA	59.89
25N	25C19	230	48.33	25N	25C19	23C	44.21
25N	25C19	162NE2	80.81	25 <b>N</b>	25C19	19CD	47.37
25N	25C19	190E1	51.79	25 <b>CA</b>	25C19	23CA	78.48
25CA	25C19	162CA	93.16	25CA	25C19	230	66.60
25CA	25C19	23C	63.45	25 <b>CA</b>	25C19	162CG	82.49

TABLE XIX									
25CA	25C19	162NE2	65.00	25CA	25C19	19CD	49.86		
25CA	25C19	190E1	47.62	25CA	25C19	162CB	95.23		
23CA	25C19	230	31.44	23CA	25C19	23C	19.68		
23CA	25C19	19CD	59.89	23CA	25C19	190E1	75.03		
162CA	25C19	162CG	32.75	162CA	25C19	162NE2	59.16		
162CA	25C19	161C	29.34	162CA	25C19	162CB	18.30		
230	25C19	23C	15.83	230	25C19	19CD	75.71		
230	25C19	190E1	88.43	23C	25C19	19CD	61.25		
23C	25C19	190E1	74.96	162CG	25C19	162NE2	27.82		
162CG	25C19	19CD	86.48	162CG	25C19	190E1	71.46		
162CG	25C19	161C	57.47	162CG	25C19	162CB	18.04		
162NE2	25C19	19CD	58.67	162NE2	25C19	190E1	43.65		
162NE2	25C19	161C	85.29	162NE2	25C19	162CB	45.84		
19CD	25C19	190E1	15.17	190E1	25C19	162CB	89.49		
161C	25C19	162CB	39.48	19NE2	25020	25 <b>S</b> G	93.68		
19NE2	25020	23CA	67.16	19NE2	25020	25CB	65.25		
19NE2	25020	23C	68.09	19NE2	25020	19CD	11.46		
19NE2	25020	230	84.29	19NE2	25020	25N	55.54		
19NE2	25020	190E1	27.77	19NE2	25020	24N	52.93		
19NE2	25020	162CD2	84.48	19NE2	25020	23N	61.60		
19NE2	25020	25CA	59.16	19NE2	25020	220	33.73		
25SG	25020	25CB	29.49	25 <b>S</b> G	25020	23C	86.92		
25SG	25020	19CD	84.98	25 <b>S</b> G	25020	230	77.98		
25 <i>S</i> G	25020	25N	47.84	25SG	25020	190E1	75.69		
25 <i>S</i> G	25020	24N	81.49	25SG	25020	162CD2	54.46		
25SG	25020	25CA	35.73	25 <b>SG</b>	25020	1610	62.50		
23CA	25020	23C	23.99	23CA	25020	19CD	77.34		
23CA	25029	230	36.67	23CA	25020	25 <b>N</b>	69.78		
23CA	25020	190E1	93.50	23CA	25020	24N	33.03		
23CA	25020	23N	10.73	23CA	25020	25CA	87.65		
23CA	25020	220	36.27	25 <b>CB</b>	25020	23C	83.77		
25CB	25020	19CD	55.78	25CB	25020	230	83.59		
25CB	25020	25N	34.83	25CB	25020	190E1	46.44		
25CB	25020	24N	71.21	25CB	25020	162CD2	47.80		
25CB	25020	25CA	16.76	25CB	25020	220	91.65		
25CB	25020	1610	85.05	23C	25020	19CD	75.06		
23C	25020	230	17.82	23C	25020	25 <b>N</b>	49.05		
23C	25020	190E1	88.41	23C	25020	24N	16.95		

•			TA	BLE XIX			
23C	25020	23N	33.23	23C	25020	25CA	67.04
23C	25020	220	47.64	19CD	25020	230	89.75
19CD	25020	25N	52.63	19CD	25020	190E1	16.53
19CD	25020	24N	58.75	19CD	25020	162CD2	73.33
19CD	25020	23N	72.54	19CD	25020	25CA	52.30
19CD	25020	220	45.08	230	25020	25N	51.85
230	25020	24N	31.38	230	25020	23N	47.25
230	25020	25CA	68.11	230	25020	220	65.37
25N	25020	190E1	55.38	25N	25020	24N	36.84
25N	25020	162CD2	82.30	25N	25020	23N	74.87
25N	25020	25CA	18.06	25N	25020	220	66.79
190E1	25020	24N	71.53	190E1	25020	162CD2	56.81
190E1	25020	23N	89.05	190E1	25020	25CA	48.64
190E1	25020	220	61.51	24N	25020	23 <b>N</b>	38.39
24N	25020	25CA	54.62	24N	25020	220	40.31
162CD2	25020	25 <b>CA</b>	64.39	162CD2	25020	1610	59.56
23N	25020	25CA	92.03	23 <b>N</b>	25020	220	28.59
25CA	25020	220	79.45	25CA	25020	1610	97.28
25 <i>S</i> G	25C21	1610	94.20	25SG	25C21	230	80.20
25SG	25C21	25CB	11.32	25SG	25C21	23C	74.97
25SG	25C21	161C	90.47	25SG	25C21	23CA	85.24
25SG	25C21	162CA	61.66	25 <b>S</b> G	25C21	25N	34.26
25SG	25C21	162CD2	42.75	1610	25C21	25CB	99.42
1610	25C21	161C	6.02	1610	25C21	162CA	32.54
1610	25C21	162CD2	63.74	230	25C21	25CB	75.76
230	25C21	23C	15.44	230	25C21	65CA	51.45
230	25C21	23CA	31.46	230	25C21	25N	46.40
25CB	25C21	23C	68.05	25CB	25C21	161C	96.56
25CB	25C21	23CA	75.96	25 <b>CB</b>	25C21	162CA	67.42
25CB	25C21	25N	29.59	25CB	25C21	162CD2	41.05
23C	25C21	65CA	65.91	23C	25C21	23CA	19.15
23C	25C21	25N	41.03	65CA	25C21	23CA	74.18
65CA	25C21	25N	90.25	161C	25C21	162CA	29.15
161C	25C21	162CD2	63.15	23 <b>CA</b>	25C21	25N	54.22
23CA	25C21	162CD2	99.59	162CA	25C21	25N	95.84
162CA	25C21	162CD2	38.23	25N	25C21	162CD2	69.67
1610	25N22	161C	14.11	1610	25N22	162CA	41.10
1610	25N22	162N	27.68	1610	25N22	163N	72.60

			TA	BLE XIX			
1610	25N22	162C	56.36	1610	25N22	161CA	21.99
25 <i>S</i> G	25N22	162CA	74.63	25 <i>S</i> G	25N22	162N	94.42
25 <b>S</b> G	25N22	163N	59.02	25 <b>S</b> G	25 <b>N2</b> 2	25CB	3.31
25 <b>S</b> G	25N22	162C	70.77	161C	25N22	162CA	35.66
161C	25N22	162N	17.48	161C	25N22	163N	63.91
161C	25N22	162C	47.46	161C	25N22	161CA	15.87
162CA	25N22	162N	19.79	162CA	25N22	163N	32.40
162CA	25N22	25CB	71.33	162CA	25N22	162C	17.88
162CA	25N22	161CA	50.91	162N	25N22	163N	46.46
162N	25N22	25CB	91.12	162N	25 <b>N22</b>	162C	29.99
162N	25N22	161CA	31.55	163N	25N22	25CB	56.10
163N	25N22	162C	16.47	163N	25 <b>N22</b>	161CA	76.35
25CB	25N22	162C	67.57	162C	25 <b>N2</b> 2	161CA	60.36
1610	25C23	25 <b>S</b> G	74.84	1610	25C23	161C	14.77
1610	25C23	162N	24.77	1610	25C23	162CA	32.09
1610	25C23	161CA	26.81	25 <b>S</b> G	25C23	161C	82.13
25SG	25C23	65CA	98.69	25SG	25C23	162N	73.79
25SG	25C23	162CA	57.15	25SG	25C23	26CD1	66.11
25SG	25C23	161CA	99.24	25SG	25C23	26CB	66.49
161C	25C23	162N	14.87	161C	25C23	162CA	29.26
161C	25C23	161CA	17.21	660	25C23	65CA	69.84
660	25C23	66 <b>N</b>	37.37	660	25C23	65C	52.20
660	25C23	26CD1	54.29	660	25C23	26CB	42.11
65CA	25C23	66N	32.68	65CA	25C23	65C	17.67
65CA	25C23	26CD1	48.54	65CA	25C23	26CB	74.99
66N	25C23	65C	15.04	6 <b>6N</b>	25C23	26CD1	43.34
66N	25C23	26CB	56.16	162N	25C23	162CA	17.00
162N	25C23	161CA	29.22	162CA	25C23	161CA	45.57
65C	25C23	26CD1	42.77	65C	25C23	26CB	63.55
26CD1	25C23	26CB	29.73	65 <b>CA</b>	25024	66N	44.96
65CA	25024	65C	24.04	65CA	25024	660	88.80
65CA	25024	65 <b>N</b>	11.32	65CA	25024	26CD1	59.03
65CA	25024	66CA	56.76	65CA	25024	66C	76.00
65CA	25024	6 <b>4</b> 0	31.19	65CA	25024	230	51.20
65CA	25024	26CG	69.83	65CA	25024	26CB	87.37
65CA	25024	650	23.39	65CA	25024	64C	20.95
66N	25024	65C	20.92	66N	25024	660	44.37
6 <b>6N</b>	25024	65N	56.05	66N	25024	26CD1	51.61

	TABLE XIX									
66N	25024	66CA	11.81	66N	25024	66C	31.20			
66N	25024	640	68.01	66N	25024	230	77.65			
66N	25024	26CG	49.90	66N	25024	26CB	61.70			
66N	25024	650	21.59	66N	25024	64C	64.08			
65C	25024	660	65.02	65C	25024	65N	35.18			
65C	25024	26CD1	51.98	65C	25024	66CA	32.73			
65C	25024	66C	52.03	65C	25024	640	49.51			
65C	25024	230	64.02	65C	25024	26CG	57.07			
65C	25024	26CB	72.88	65C	25024	650	0.87			
65C	25024	64C	43.58	660	25024	65N	99.41			
660	25024	26CD1	61.59	660	25024	66CA	32.76			
660	25024	66C	13.43	660	25024	26CG	47.99			
660	25024	26CB	43.51	660	25024	650	65.76			
65N	25024	26CD1	62.12	65N	25024	66CA	67.82			
65N	25024	66C	86.89	65N	25024	640	27.80			
65N	25024	230	45.06	65N	25024	26CG	74.86			
65N	25024	26CB	92.41	65N	25024	650	34.57			
65N	25024	64C	14.20	26CD1	25024	25SG	67.04			
26CD1	25024	66CA	53.75	26CD1	25024	66C	58.39			
26CD1	25024	640	89.45	26CD1	25024	230	41.01			
26CD1	25024	26CG	15.28	26CD1	25024	26CB	31.14			
26CD1	25024	650	52.64	26CD1	25024	64C	76.32			
25 <b>S</b> G	25024	1610	59.11	25 <b>S</b> G	25024	230	57.25			
25 <b>S</b> G	25024	26CG	<b>69</b> .72	25SG	25024	26CB	63.45			
66CA	25024	66C	19.46	66CA	25024	640	78.88			
66CA	25024	230	85.60	66CA	25024	26CG	48.11			
66CA	25024	26CB	56.41	66CA	25024	650	33.40			
66CA	25024	64C	75.76	66C	25024	640	97.72			
66C	25024	230	96.86	66C	25024	26CG	47.34			
66C	25024	26CB	48.38	6 <b>6C</b>	25024	650	52.74			
6 <b>6C</b>	25024	64C	95.21	640	25024	230	69.05			
640	25024	650	48.56	640	25024	64C	14.27			
230	25024	26CG	55.76	230	25024	26CB	67.79			
230	25024	650	64.06	230	25024	64C	55.23			
26CG	25024	26CB	17.62	26CG	25024	650	57.87			
26CG	25024	64C	88.94	26CB	25024	650	73.72			
650	25024	64C	42.81	1610	25C25	161C	18.10			
1610	25C25	162N	30.50	1610	25C25	161CA	32.52			

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		TA	BLE XIX		
1610	25C25 163N	64.43	1610	25C25 25SG	63.34
1610	25C25 162CA	35.10	1610	25C25 162C	53.40
1610	25C25 163CB	93.18	660	25C25 25SG	99.66
660	25C25 66N	34.44	660	25C25 163CB	83.50
161C	25C25 162N	17.33	161C	25C25 161CA	19.80
161C	25C25 163N	60.64	161C	25C25 25SG	75.44
161C	25C25 162CA	30.92	161C	25C25 162C	46.09
161C	25C25 163CB	89.51	162N	25C25 161CA	32.24
162N	25C25 163N	44.92	162N	25C25 25SG	71.34
162N	25C25 162CA	18.08	162N	25C25 162C	29.44
162N	25C25 163CB	73.04	161CA	25C25 163N	77.08
161CA	25C25 25SG	94.58	161CA	25C25 162CA	49.19
161CA	25C25 162C	61.23	163N	25C25 25SG	49.61
163N	25C25 162CA	30.25	163N	25C25 162C	16.24
163N	25C25 163CB	29.19	25SG	25C25 162CA	54.70
25SG	25C25 162C	58.88	25 <b>S</b> G	25C25 66N	88.50
25SG	25C25 163CB	57.12	162CA	25C25 162C	18.49
162CA	25C25 163CB	59.44	162C	25C25 163CB	43.61
660	25C26 26CB	56.41	660	25C26 66C	2.37
660	25C26 26CA	74.83	660	25C26 68SD	74.84
660	25C26 66N	33.64	660	25C26 26N	86.28
660	25C26 26CG	50.23	660	25C26 26CD1	56.63
163CB	25C26 26CB	71.13	163CB	25C26 163N	35.96
163CB	25C26 25SG	68.31	163CB	25C26 163CA	18.75
163CB	25C26 26CA	53.74	163CB	25C26 162C	49.97
163CB	25C26 68SD	50.69	163CB	25C26 1610	96.76
163CB	25C26 26N	55.09	163CB	25C26 26CG	85.31
163CB	25C26 162CA	65.37	163CB	25C26 161C	92.67
163CB	25C26 162N	78.07	163CB	25C26 26CD1	92.73
26CB	25C26 163N	98.44	26CB		73.00
26CB	25C26 163CA	88.50		25C26 66C	55.68
26CB	25C26 26CA	18.86	26CB	25C26 68SD	60.85
26CB	25C26 66N	60.69	25CB	25C26 26N	31.12
26CB	25C26 26CG	15.82	26CB	25C26 26CD1	29.79
163N	25C26 25SG	54.34	163ท	25C26 163CA	19.23
163N	25C26 25CA	79.60	163N	25C26 162C	16.17
163N	25C26 68SD	84.27	163N	25C26 1610	61.07
163N	25C26 26N	71.27	163N	25C26 162CA	29.41

			TA	BLE XIX			
163N	25C26	161C	57.02	163N	25C26	162N	43.18
25 <i>S</i> G	25C26	163CA	65.77	25SG	25C26	26CA	61.96
25 <b>S</b> G	25C26	162C	61.73	25 <i>S</i> G	25C26	1610	58.28
25SG	25C26	66N	92.45	25SG	25C26	26N	44.25
25 <i>S</i> G	25C26	26CG	71.07	25 <i>S</i> G	25C26	162CA	54.19
25 <i>S</i> G	25C26	161C	68.75	25 <b>S</b> G	25C26	162N	67.87
25SG	25C26	26CD1	61.51	163CA	25C26	26CA	70.22
163CA	25C26	162C	31.46	163CA	25C26	68SD	65.05
163CA	25C26	1610	79.96	163CA	25C26	26N	67.43
163CA	25C26	162CA	47.86	163CA	25C26	161C	74.37
163CA	25C26	162N	59.50	66C	25C26	26CA	74.25
66C	25C26	68SD	76.55	66C	25C26	6 <b>6N</b>	31.36
66C	25C26	26N	85.17	66C	25C26	26CG	48.83
66C	25C26	26CD1	54.70	26CA	25C26	162C	95.72
26CA	25C26	68SD	58.01	26CA	25C26	6 <b>6N</b>	77.90
26CA	25C26	26N	17.81	26CA	25C26	26CG	31.59
26CA	25C26	26CD1	40.72	162C	25C26	68 <i>S</i> D	93.90
162C	25C26	1610	49.88	162C	25C26	26N	86.44
162C	25C26	162CA	18.09	162C	25C26	161C	42.92
162C	25C26	162N	28.10	68SD	25C26	26N	73.74
68SD	25C26	26CG	75.07	68SD	25C26	26CD1	90.33
1610	25C26	162CA	32.21	1610	25C26	161C	14.49
1610	25C26	162N	26.44	66N	25C26	26N	79.75
66N	25C26	26CG	46.31	6 6 N	25C26	26CD1	41.21
26N	25C26	26CG	36.93	26N	25C26	162CA	89.41
26N	25C26	26CD1	38.63	26CG	25C26	26CD1	15.55
162CA	25C26	161C	28.03	162CA	25C <b>26</b>	162N	16.67
161C	25C26	162N	15.37	163CB	25C27	68SD	63.80
163CB	25C27	163CA	20.61	163CB	25C27	163N	36.17
163CB	25C <b>27</b>	68CE	66.45	163CB	25C27	134CB	82.27
163CB	25C27	26CB	68.13	163CB	25C27	162C	50.98
163CB	25C27	1620	55.96	163CB	25C27	26CA	50.39
68SD	25C27	660	85.83	69 <b>S</b> D	25C27	163CA	78.97
68SD	25C27	163N	98.79	68SD	25C27	68CE	27.50
68SD	25C27	134CB	86.61	68SD	25C27	209CD2	77.07
63SD	25C27	26CB	65.47	68SD	25C27	66C	82.65
68SD	25C27	26CA	60.55	68SD	25C27	67CA	62.19
660	25C <b>27</b>	26CB	49.09	660	25C27	66C	4.22

			TAI	BLE XIX			
660	25C27	26CA	66.81	660	25C27	67CA	32.70
163CA	25C27 1	L63N	20.52	163CA	25C27	68CE	72.92
163CA	25C27 1	L34CB	68.04	163CA	25C27	26CB	86.68
163CA	25C27 1	L62C	32.08	163CA	25C27	1620	35.38
163CA	25C27	26CA	69.20	163N	25C27	68CE	93.00
163N	25C27 1	L34CB	74.40	163N	25C27	26CB	90.05
163N	25C27 1	L62C	16.34	163N	25C27	1620	27.46
163N	25C27	26CA	74.74	68CE	25C27	134CB	59.27
68CE	25C27 2	209CD2	56.93	68CE	25C27	26CB	92.20
68CE	25C27 1	L62C	98.29	68CE	25C27	1620	89.77
68CE	25C27	26CA	84.17	68CE	25C27	67CA	79.42
134CB	25C27 2	209CD2	45.82.	134CB	25C27	162C	65.01
134CB	25C27 1	L620	50.58	209CD2	25C27	1620	95.82
209CD2	25C27	67CA	79.47	26CB	25C27	66C	50.07
26CB	25C27	26CA	17.74	26CB	25C27	67CA	62.22
162C	25C27 1	1620	14.47	162C	25C27	26CA	90.70
66C	25C27	26CA	67.64	66C	25C27	67CA	28.48
26CA	25C27	67CA	76.24	134CB	25C28	163CB	94.94
134CB	25C28 1	163N	95.08	134CB	25C28	162C	86.92
134CB	25C28 1	1620	68.97	134CB	25C28	163CA	83.27
134CB	25C28 2	209CD2	54.77	134CB	25C28	134CA	19.96
134CB	25C28 1	162N	96.63	134CB	25C28	68CE	62.89
134CB	25C28	68SD	87.41	134CB	25C28	134C	23.58
134CB	25C28 1	161N	87.00	163CB	25C28	163N	38.06
163CB	25C28 1	162C	57.44	163CB	25C28	1620	66.17
163CB	25C28	163CA	22.67	163CB	25C28	134CA	78.61
163CB	25C28	162N	88.44	163CB	25C28	162CA	69.97
163CB	25C28	68CE	60.09	163CB	25C28	68SD	51.61
163CB	25C28	161C	98.14	163CB	25C28	660	84.86
163CB	25C28	134C	88.60	163CB	25C28		91.88
163N	25C28	162C	20.41	163N	25C28	1620	35.09
163N	25C28	163CA	22.12	163N	25C28	134CA	75.15
163N	25C28	162N	50.53	163N	25C28	162CA	32.37
163N	25C28	68CE	93.09	163N	25C28	68SD	89.55
163N	25C28	161C	62.08	163N	25C28		76.23
163N	25C28	1610	59.80	163N	25C28		83.44
162C	25C28	1620	18.65	162C		163CA	37.43
162C	25C28	134CA	68.48	162C	25C28	162N	33.45

TABLE XIX										
162C	25C28	162CA	18.81	162C	25C28	161C	48.13			
162C	25C28	134C	64.60	162C	25C28	1610	50.86			
162C	25C28	161N	64.31	1620	25C28	163CA	43.61			
1620	25C28	134CA	51.60	1620	25C28	162N	38.24			
1620	25C28	162CA	32.12	1620	25C28	161C	54.89			
1620	25C28	134C	46.14	1620	25C28	1610	62.22			
1620	25C28	161N	60.29	163CA	25C28	134CA	64.10			
163CA	25C28	162N	70.64	163CA	25C28	162CA	53.50			
163CA	25C28	68CE	71.06	163CA	25C28	68SD	70.42			
163CA	25C28	161C	83.69	163CA	25C28	134C	70.58			
163CA	25C28	1610	81.89	209CD2	25C28	134CA	73.08			
209CD2	25C28	68CE	56.44	209CD2	25C28	68SD	70.67			
209CD2	25C28	660	90.11	209CD2	25C28	134C	77.7 <b>7</b>			
134CA	25C28	162N	84.78	134CA	25C28	162CA	83.66			
134CA	25C28	68CE	63.87	134CA	25C28	68SD	86.25			
134CA	25C28	161C	99.94	134CA	25C28	134C	14.94			
134CA	25C28	161N	85.92	162N	25C28	162CA	18.62			
162N	25C28	161C	16.67	162N	25C28	134C	73.76			
162N	25C28	1610	26.48	162N	25C28	161N	33.69			
162CA	25C28	161C	30.27	162CA	25C28	134C	76.50			
162CA	25C28	1610	32.09	162CA	25C28	161N	52.27			
68CE	25C28	68SD	24.56	68CE	25C28	660	83.70			
68CE	25C28	134C	78.45	68SD	25C28	660	61.76			
161C	25C28	134C	87.61	161C	25C28	1610	14.08			
161C	25C28	161N	30.20	660	25C28	1610	96.95			
134C	25C28	151N	71.12	1610	25C28	161N	43.13			
660	25C29	68SD	95.60	650	25C29	67CA	46.15			
660	25C29	67CD1	76.62	660	25C29	66C	11.19			
660	25C29	67N	28.20	660	25C29	68N	68.80			
660	25C29	67C	53.21	660	25C29	67CB	59.40			
660	25C29	67CG	63.57	560	25C29	234OH2	97.74			
660	25C29	26CB	45.24	660	25C29	67CE1	81.98			
209CD2	25C29	68SD	86.14	209CD2	25C29	67CD1	76.20			
209CD2	25C29	68CE	61.12	209CD2	25C29	68 <b>N</b>	93.85			
209CD2	25C29	67CB	92.20	209CD2	25C29	67CG	87.96			
209CD2	25C29	134CB	44.39	209CD2	25C29	234OH2	59.00			
209CD2	25C29	67CE1	74.42	683D	25C29	67CA	76.30			
68SD	25C29	66C	94.73	68SD	25029	68CE	26.92			

			TAI	BLE XIX			
68SD	25C29	67N	86. <b>9</b> 0	68SD	25C29	68N	44.21
68SD	25C29	67C	58.54	68SD	25C29	67CB	85.01
68SD	25C29	163CB	50.89	68SD	25C29	134CB	79.09
68SD	25C29	2340H2	57.89	68SD	25C29	26CB	61.09
67CA	25C29	67CD1	50.51	67CA	25C29	66C	35.89
67CA	25C29	68CE	93.88	67CA	25C29	67N	18.53
67CA	25C29	68N	33.04	67CA	25C29	67C	17.87
67CA	25C29	67CB	18.28	67CA	25C29	67CG	34.89
67CA	25C29	234OH2	51.94	67CA	25C29	26CB	68.39
67CA	25C29	67CE1	64.72	67CD1	25C29	66C	66.65
67CD1	25C29	67N	56.05	67CD1	25C29	68N	72.69
67CD1	25C29	67C	65.26	67CD1	25C29	67CB	33.63
67CD1	25C29	67CG	15.94	67CD1	25C29	2340H2	57.55
67CD1	25C29	67CE1	15.80	66C	25C29	67N	17.49
66C	25C29	68N	61.93	66C	25C29	67C	45.45
66C	25C29	67CB	48.27	66C	25C29	67CG	52.89
66C	25C29	2340H2	87.82	66C	25C29	26CB	52.23
66C	25C29	67CE1	73.67	68CE	25C29	68N	61.24
68CE	25C29	67C	77.92	68CE	25C29	67CB	95.69
68CE	25C29	163CB	55.77	68CE	25C29	134CB	54.85
68CE	25C29	2340H2	55.46	68CE	25C29	26CB	86.35
67N	25C29	68N	47.83	67N	25C29	67C	30.97
67N	25C29	67CB	31.65	67N	25C29	67CG	40.58
67N	25C29	2340H2	70.44	67 <b>N</b>	25C29	26CB	60.08
67N	25C29	67CE1	66.83	6 <b>8N</b>	25C29	67C	16.87
68N	25C29	67CB	41.26	68N	25C29	67CG	60.50
68N	25C29	163CB	90.93	6 <b>8N</b>	25C29	2340H2	36.37
68N	25C29	26CB	64.40	6 <b>8N</b>	25C29	67CE1	88.49
67C	25C29	67CB	31.65	67C	25C29	67CG	50.66
67C	25C29	163CB	99.57	67C	25C29	2340H2	46.71
67C	25C29	26CB	59.86	67C	25C29	67CE1	80.48
67CB	25C29	67CG	19.43	67CB	25C29		44.01
67CB	25C29	26CB	86.67	67CB	25C29	67CE1	48.89
67CG	25C29	2340H2	54.79	67 <b>CG</b>	25C29	67CE1	29.95
163CB	25C29	134CB	64.77	163CB	25C29	26CB	55.99
134CB	25C29	2340H2	92.26	2340H2	25C29	67CE1	70.61
660	25N30	66N	36.00	660	25N30	66C	10.42
66N	25N30	66C	31.15	1.610	25N30	161C	14.88

TABLE XIX									
1610	25N30	161CA	29.27	1610	25N30	1600	61.86		
161C	25N30	161CA	18.36	161C	25N30	1600	49.77		
161CA	25N30	1600	32.60	1600	25C31	161CA	38.22		
1600	25C31	160C	13.90	1600	25C31	161C	56.07		
1600	25C31	161N	27.07	1600	25C31	1610	68.03		
161CA	25C31	160C	31.81	161CA	25C31	161C	19.14		
161CA	25C31	161N	17.82	161CA	25C31	1610	29.81		
160C	25C31	161C	46.85	160C	25C31	161N	16.04		
160C	25C31	1610	60.44	161C	25C31	161N	30.84		
161C	25C31	1610	14.70	660	25C31	67CE1	65.45		
161N	25C31	1610	44.68	1600	25032	160C	20.78		
1600	25032	161CA	51.99	1600	25032	161N	38.37		
1600	25032	161C	74.30	1600	25032	160CB	45.78		
1600	25032	1610	86.34	1600	25032	160CA	28.59		
1600	25032	162N	77.29	1600	25032	161CB	49.03		
160C	25032	161CA	42.82	160C	25032	161N	21.72		
160C	25032	161C	61.05	160C	25032	160CB	36.54		
160C	25032	1610	75.94	160C	25032	160CA	16.12		
160C	25032	162N	59.68	160C	25032	161CB	45.43		
161CA	25032	161N	24.13	161CA	25032	161C	23.46		
161CA	25032	160CB	75.84	161CA	25032	1610	34.47		
161CA	25032	160CA	57.38	161CA	25032	162N	33.92		
161CA	25032	161CB	11.93	161N	25032	161C	39.41		
161N	25032	160CB	51.73	161N	25032	1610	54.76		
161N	25032	160CA	34.01	161N	25032	162N	39.00		
161N	25032	161CB	31.34	161C	25032	160CB	86.66		
161C	25032	1610	16.47	161C	25032	160CA	72.47		
161C	25032	162N	16.93	161C		161CB	32.27		
160CB	25032	160CA	20.46	160CB	25032		76.70		
160CB	25032	161CB	81.15	1610		160CA	88.45		
1610	25032	162N	2 <b>9</b> .70	1610	25032	161CB	38.78		
160CA	25032	162N	66.95	160CA	25032	161CB	61.27		
162N	25032	161CB	45.15	67CE1	25C33	67CZ	16.24		
67CE1	25C33	67CD1	18.33	67CE1	25C33	670H	30.78		
67CZ	25C33	67CD1	31.64	67CZ	25 <b>C</b> 33		17.77		
67CD1	25C33	67 <b>0H</b>	48.37	1600	25C34	160C	9.85		
1600	25C34	160CB	38.05	67CE1	25C34	670H	31.14		
67CE1	25C34	67CZ	15.25	67CE1	25C34	67CD1	15.11		

	TABLE XIX									
670H	25C34	67CZ	17.26	670H	25C34	67CD1	45.96			
67CZ	25C34	67CD1	29.18	160C	25C34	160CB	29.90			
67CE1	25C35	670H	32.70	67CE1	25C35	209CD2	77.34			
67CE1	25C35	67CZ	14.99	67CE1	25C35	67CD1	16.08			
67CE1	25C35	209CD1	77.10	670H	25C35	67CZ	17.98			
670H	25C35	67CD1	48.74	670H	25C35	209CD1	99.59			
209CD2	25C35	67CZ	92.22	209CD2	25C35	67CD1	62.32			
209CD2	25C35	1600	97.37	209CD2	25C35	160CB	70.82			
209CD2	25C35	209CD1	31.32	67CZ	25C35	67CD1	30.87			
67CZ	25C35	209CD1	88.81	67CD1	25C35	209CD1	67.63			
1600	25C35	160CB	37.56	160CB	25C35	209CD1	83.17			
1600	25N36	160CB	39.26	1600	25N36	160CD1	70.13			
1600	25N36	160CG	56.11	160CB	25N36	160CD1	30.89			
160CB	25N36	160CG	18.76	160CD1	25N36	160CG	18.07			
67CE1	25N36	670H	27.00	1600	25C37	160CB	42.20			
1600	25C37	160N	37.40	1600	25C37	160C	10.47			
1600	25C37	160CA	28.50	1600	25C37	160CG	58.56			
1600	25C37	1580	81.42	160CB	25C37	160N	31.80			
160CB	25C37	160C	31.91	160CB	25C37	160CA	18.43			
160CB	25C37	160CG	17.89	160CB	25C37	1580	70.66			
160N	25C37	160C	30.12	160N	25C37	160CA	17.61			
160N	25C37	160CG	36.16	160N	25C37	1580	47.06			
160C	25C37	160CA	18.39	160C	25C37	160CG	48.09			
160C	25C37	1580	76.49	160CA	25C37	160CG	31.12			
160CA	25C37	1580	63.78	160CG	25C37	1580	60.96			
1600	25N38	160C	7.90	1600	25N38	160CB	39.77			
1600	25N38	160N	32.65	1600	25N38	160CA	24.13			
160C	25N38	160CB	32.02	160C	25N38	160N	29.52			
160C	25N38	160CA	17.57	160CB	25N38	160N	29.89			
160CB	25N38	160CA	18.19	160N	25N38	160CA	17.29			

#### TABLE XX

Table of distances in Angstroms between atoms of the inhibitor and protein for all protein atoms within 5 Angstroms of the inhibitor 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	2420H2	3.097	25C1	180D1	3.745	25C1	184CB	3.754
25C1	184CA	3.830	25C1	18ND2	3.916	25C1	1840	3.932
25C1	184CG	3.975	25C1	184CD1	4.022	25C1	18CG	4.158
25C1	184C	4.165	25C1	184CD2	4.830	25C1	184NE1	4.840
25C1	200	4.895	25C2	180D1	2.946	25C2	184CD1	3.510
25C2	184CA	3.592	25C2	18CG	3.620	25C2	20N	3.785
25C2	18ND2	3.842	25C2	184CB	3.886	25C2	2420H2	3.894
25C2	184CG	3.897	25C2	200	3.948	25C2	1830	4.106
25C2	184C	4.198	25C2	19CG	4.207	25C2	20CA	4.229
25C2	19N	4.334	25C2	.1840	4.367	25C2	184NE1	4.377
25C2	20C	4.510	25C2	184N	4.678	25C2	19C	4.809
25C2	18CB	4.827	25C2	183C	4.828	25C2	18CA	4.864
25C2	18C	4.870	25C2	184CD2	4.940	25C2	19CA	4.979
25C3	200	2.991	25C3	184CD1	3.420	25C3	19CG	3.460
25C3	20N	3.592	25C3	180D1	3.738	25C3	20C	3.852
25C3	184NE1	3.896	25C3	20CA	4.002	25C3	184CG	4.180
25C3	19CD	4.295	25C3	19C	4.443	25C3	19N	4.455
25C3	184CA	4.500	25C3	183C	4.509	25C3	19CB	4.596
25C3	184CB	4.635	25C3	18CG	4.662	25C3	190E1	4.690
25C3	19CA	4.707	25C3	184CE2	4.837	25C3	2420H2	4.986
25C4	200	3.250	25C4	184CD1	3.862	25C4	184NE1	3.932
25C4	20C	4.284	25C4	19CG	4.378	25C4	184CG	4.519
25C4	184CE2	4.630	25C4	20N	4.702	25 <b>C4</b>	20CA	4.823
25C4	19CD	4.851	25C4	21NE2	4.914	25C4	180D1	4.958
25C4	184CD2	4.971	25C5	134CD1	4.323	25C5	200	4.342
25C5	184NE1	4.439	25C5	1.84CG	4.577	25C5	184CE2	4.756
25C5	2420H2	4.835	25C5	184CD2	4.858	25C5	21NE2	4.951
25C6	2420H2	3.703	25C6	184CG	4.325	2506	184CD1	4.403

# TABLE XX

25C6	184CB	4.414	25C6	184CD2	4.792	25C6	184NE1	4.871
25C6	184CA	4.873	25C6	180D1	4.955	25C6	1840	4.967
25C7	200	3.169	25C7	184NE1	4.191	25C7	20C	4.372
25C7	19CG	4.412	25C7	19CD	4.508	25C7	184CD1	4.544
25C7	19NE2	4.715	25C7	190E1	4.899	25C7	184CE2	4.927
2508	184NE1	3.443	2508	19CD	3.459	2508	19NE2	3.624
2508	190E1	3.712	2508	19CG	3.802	2508	200	3.894
2508	184CD1	4.085	2508	184CE2	4.312	2508	220	4.666
2508	184CZ2	4.747	25C9	19NE2	3.833	25C9	184NE1	3.925
25C9	19CD	3.963	25C9	190E1	4.023	25C9	184CE2	4.572
25C9	184CZ2	4.604	25C9	19CG	4.740	25C9	184CD1	4.906
25C9	220	4.956	25010	19NE2	3.975	25010	23CA	4.264
25010	19CD	4.483	25010	220	4.554	25010	23N	4.718
25010	190E1	4.724	25010	22C	4.857	25C11	162ND1	3.919
25C11	184CZ2	3.933	25C11	162CE1	4.435	25C11	184NE1	4.520
25C11	184CE2	4.600	25C11	162CG	4.656	25C11	184CH2	4.932
25C11	162CB	4.962	25C13	1380G	4.932	25C15		3.572
25C15	138CB	4.696	25C15	138CA	4.824	25C15	1610D1	4.950
25C16	162ND1	3.171	25C16	1610	3.822	25C16		3.975
25C16	162CE1	4.007	25C16	162CB	4.112	25C16	25SG	4.187
25C16	162CA	4.258	25C16	161C	4.677		184CZ2	4.751
25C16	25CB	4.882	25C16	162N	4.893	25C16	•	4.906
25017	162ND1	2.637	25017	162CB	2.951	25017		3.023
25017	162CG	3.092	25017	162CA	3.124	25017		3.669
25017	162N	3.731	-25017	162CE1	3.755	25017		4.071
25017	1610D1	4.274	25017	162CD2	4.298	25017		4.467
25017	162NE2	4.594	25017	184CZ2	4.770	25017		4.786
25017	25CB	4.931	25017	161CA	4.957	25017		4.964
25N18	25 <b>S</b> G	3.671		162ND1	3.834	25N18		3.966
25N18	25CB	4.503		19NE2			23CA	4.510
25N18	162CE1	4.582	25N18	162CA	4.791			
25N18	162CG	4.836		162CB				
	1610			162NDI				
25C19	162CA	4.350	25C19					
25C19	230	4.704	25C19	23C	4.758		162N	
25C19	25N	4.315	25019	19NE2	4.957	25C19	163N	4.992

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### TABLE XX

25C19	162CE1	4.994	25C19	162CB	4.997	25C20	1610	3.580
25C20	25 <b>S</b> G	4.069	25C20	23CA	4.393	25C20	230	4.523
25C20	23C	4.786	25C20	161C	4.808	25C21	1610	3.316
25C21	161C	4.462	25C22	1610	2.892	25C22	161C	3.848
25C22	161CA	4.121	25C22	161CB	4.359	25C23	640	4.590
25C23	1610	4.811	25N24	184NE1	3.412	25N24	184CZ2	3.492
25N24	184CE2	3.750	25N24	190E1	4.106	25N24	162ND1	4.147
25N24	162CE1	4.221	25N24	19CD	4.432	25N24	19NE2	4.515
25N24	184CD1	4.633	25N24	184CH2	4.684	25C25	25SG	1.822
25C25	25CB	3.054	25C25	25 <b>N</b>	3.471	25C25	230	3.687
25C25	25CA	3.792	25C25	23C	3.793	25C25	23CA	3.994
25C25	19NE2	4.214	25C25	162ND1	4.271	25C25	1610	4.312
25C25	24N	4.350	25C25	26N	4.445	25C25	25C	4.500
25C25	24C	4.578	25C25	163N	4.693	25C25	190E1	4.797
25C25	162CA	4.833	25C25	162CE1	4.899	25C25	24CA	4.928
25C25	19CD	4.983	25026	25 <b>S</b> G	2.496	25026	25N	2.842
25026	23C	2.922	25026	25CB	2.985	25026	23CA	3.053
25026	19NE2	3.071	25026	230	3.183	25026	24N	3.337
25026	25CA	3.477	25026	24C	3.898	25026	19CD	3.997
25026	190E1	4.054	25026	24CA	4.110	25026	23N	4.413
25026	220	4.471	25026	25C	4.502	25026	26N	4.555
25026	162ND1	4.591	25026	22C	4.912	25026	162CE1	4.933
25C27	25 <b>SG</b>	2.646	25C27	230	3.049	25C27	23C	3.646
25C27	25N	3.830	25C27	65CA	3.894	25C27	25CB	4.015
25C27	26CD1	4.040	25C27	26N	4.152	25C27	23CA	4.158
25C27	25CA	4.348	25C27	24N	4.411	25C27	24C	4.634
25C27	25C	4.654	25C27	1610	4.714	25C27	65N	4.725
25C27	26CB	4.746	25C27	66N	4.758	25C27	24CA	4.780
25C27	26CG	4.815	25C27	65C	4.906	25028	25SG	3.234
25028	65CA	3.931	25028	1610	3.970	25028	230	4.277
25028	6 <b>6N</b>	4.333	25028	660	4.402	25028	26CD1	4.553
25028	65C	4.712	25028	25N	4.830	25028	26CB	4.858
25028	161C	4.903	25028	25CB	4.910	25028	163N	4.929
25C29	660	3.071	25C29	66N	3.131	25C29	65CA	3.415
25C29	65C	3.768	25C29	66C	4.000	25C29	26CD1	4.084
25C29	66CA	4.184	25C29	26CB	4.278	25C29	25SG	4.306

# TABLE XX

25C29	26CG	4.529	25C29	230	4.746	25C29	65N	4.756
25C29	1610	4.948	25C29	650	4.978	25C29	26N	4.985
25C30	660	3.147	25C30	66N	3.696	25C30	66C	4.040
25C30	65CA	4.366	25C30	66CA	4.428	25C30	65C	4.461
25C30	1610	4.799	25C31	660	3.488	25C31	1610	4.535
25C31	66C	4.544	25C31	161C	4.739	25C31	66N	4.865
25031	163CB	4.886	25C31	1600	4.980			

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#### TABLE XXI

Table of distances in Ångstroms between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom 1	Atom 2	Dist.
25C1	184CB	3.966	25C1	184CG.	4.092	25C1	180D1	4.178
25C1	184CD1	4.335	25C1	184CA	4.489	25C1	184CD2	4.689
25C1	1840	4.748	25C1	184C	4.857	25C2	180D1	3.023
25C2	184CD1	4.012	25C2	184CB	4.046	25C2	184CA	4.130
25C2	184CG	4.135	25C2	18CG	4.154	25C2	200	4.291
25C2	20N	4.526	25C2	184C	4.608	25C2	18ND2	4.696
25C2	20CA	4.804	25C2	184NE1	4.825	25C2	1840	4.835
25C2	20C	4.924	25C2	19CG	4.976	25C3	200	3.073
25C3	180D1	3.568	25C3	20C	3.908	25C3	20N	3.934
25C3	184CD1	3.954	25C3	20CA	4.129	25C3	19CG	4.226
25C3	184CG	4.484	25C3	184NE1	4.490	25C3	18CG	4.696
25C3	184CB	4.795	25C3	184CA	4.856	25C3	19C	4.885
25C3	19CD	4.898	25C4	200	3.345	25C4	184CD1	4.222
25C4	184NE1	4.344	25C4	20C	4.381	25C4	184CG	4.772
25C4	19CG	4.838	25C4	184CE2	4.948	25C4	180D1	4.956
25C4	20N	4.984	25C5	184CD1	4.521	25C5	184NE1	4.550
25C5	200	4.685	25C5	184CG	4.727	25C5	184CE2	4.764
25C5	184CD2	4.873	25C6	184CG	4.403	25C6	184CD1	4.580
25C6	184CD2	4.609	25C6	184CB	4.657	25C6	184NE1	4.884
25C6	184CE2	4.906	25C7	200	2.849	25C7	20C	4.026
25C7	184NE1	4.648	25C7	21CA	4.664	25C7	19CD	4.697
25C7	19CG	4.714	25C7	210E1	4.738	25C7	21N	4.799
25C7	184CD1	4.844	25C7	190E1	4.907	25C7	19NE2	4.961
2508	200	3.212	2508	19CD	3.453	2508	190E1	3.622
2508	19NE2	3.664	2508	19CG	3.806	2508	184NE1	3.870
2508	184CD1	4.265	2508	20C	4.432	2508	220	4.540
2508	184CE2	4.808	2508	19CB	4.983	25C9	190E1	3.889
25C9	19NE2	3.915	25 <b>C</b> 9	19CD	3.947	25C9	184NE1	4.155
25C9	200	4.391	25C9	19CG	4.723	25C9	220	4.796
25C9	184CE2	4.909	25C9	184CD1	4.929	25010	19NE2	4.578

# TABLE XXI

25010	200	4.660	25010	220	4.792	25010	19CD	4.890
25010	190E1	4.963	25C11	162ND1	4.101	25C11	184CZ2	4.175
25C11	184NE1	4.284	25C11	162CE1	4.319	25C11	190E1	4.351
25C11	184CE2	4.593	25C11	162CG	4.929	25C11	19NE2	4.945
25C11	19CD	4.953	25C15	184CZ2	4.332	25C15	184CH2	4.756
25C15	1430E1	4.901	25C15	1370	4.938	25C16	162ND1	3.174
25C16	162CE1	3.757	25C16	162CG	4.082	25C16	25SG	4.189
25C16	1610	4.344	25C16	190E1	4.427	25C16	162CB	4.457
25C16	162CA	4.770	25C16	162NE2	4.792	25C16	19NE2	4.868
25C16	184CZ2	4.883	25C16	25CB	4.955	25C16	184NE1	4.957
25C16	162CD2	4.977	25017	162ND1	2.574	25017	162CG	3.155
25017	162CB	3.298	25017	162CE1	3.479	25017	1610	3.597
25017	162CA	3.711	25017	25SG	4.141	25017	161C	4.187
25017	162CD2	4.194	25017	162N	4.282	25017	162NE2	4.329
25017	1610D1	4.530	25017	184CZ2	4.752	25017	137CB	4.842
25N18	25SG	3.590	25N18	162ND1	3.689	25N18	162CE1	4.248
25N18	1610	4.291	25N18	19NE2	4.292	25N18	190E1	4.452
25N18	25CB	4.467	25N18	23CA	4.557	25N18	162CG	4.791
25N18	19CD	4.816	25C19	25SG	2.732	25C19	1610	3.385
25C19	162ND1	3.772	25C19	25CB	4.081	25C19	162CA	4.451
25C19	161C	4.513	25C19	162CE1	4.560	25C19	23CA	4.589
25C19	25N	4.775	25C19	23C	4.776	25C19	230	4.793
25C19	162CG	4.797	25C19	19NE2	4.876	25C19	162CB	4.935
25C19	162N	4.994	25N20	184NE1	3.387	25N20	190E1	3.405
25N20	19CD	3.866	25N20	184CE2	4.004	25N20	19NE2	4.021
25N20	184CZ2	4.036	25N20	162CE1	4.357	25N20	184CD1	4.441
25 <b>N</b> 20	162ND1	4.565	25N20	19CG	4.876	25C21	25 <b>S</b> G	1.768
25C21	25CB	2.961	25C21	25N	3.320	25C21	25CA	3.665
25C21	230	3.874	25C21	23C	3.876	25C21	162ND1	3.978
25C21	23CA	4.127	25C21	19NE2	4.215	25C21	26N	4.241
25C21	24N	4.308	25C21	1610	4.319	25C21	25C	4.400
25C21	24C	4.472	25C21	162CE1	4.510	25C21	190E1	4.818
25C21	24CA	4.852	25C21	163N	4.871	25C21	162CA	4.898
25C21	19CD	4.974	25C21	25CD1	4.991	25022	25 <b>S</b> G	2.461
25022	25N	2.747	25022	25CB	2.888	25022	19NE2	3.089
25022	23C	3.118	25022	23CA	3.281	25022	25CA	3.369
25022	24N	3.372	25022	230	3.479	25022	24C	3.834

# TABLE XXI

25022	19CD	4.008	25022	190E1	4.090	25022	24CA	4.106
25022	220	4.311	25022	162ND1	4.396	25022	25C	4.432
25022	26N	4.446	25022	23N	4.563	25022	162CE1	4.605
25022	22C	4.909	25022	240	4.940	25C23	1600	4.313
25C23	670H	4.425	25C23	67CE1	4.574	25C23	160CB	4.873
25C24	1600	3.278	25C24	160CB	4.082	25C24	160C	4.283
25C24	67CE1	4.352	25C24	209CD2	4.555	25C24	160CA	4.652
25C24	670H	4.750	25C24	160N	4.847	25C25	1600	3.120
25C25	160C ·	4.287	25C25	67CE1	4.346	25C25	670H	4.617
25C25	160CB	4.802	25C25	67CZ	4.867	25C26	1600	4.072
25C26	670H	4.149	25C26	67CE1	4.572	25C26	67CZ	4.684
25C27	670H	3.768	25C27	67CZ	4.663	25C27	67CE1	4.776
25C27	1600	4.939	25C28	670H	3.920	25C28	67CE1	4.781
25C28	67CZ	4.836	25C29	1600	4.671	25C29	670H	4.760
25030	2750H2	4.167	25030	1600	4.205	25030	161CA	4.860
25C31	1600	3.769	25C31	161CA	4.036	25C31	1610	4.057
25C31	2750H2	4.224	25C31	161C	4.371	25C31	160C	4.689
25C31	161N	4.855	25C31	161CB	4.868	25032	1600	2.767
25032	161CA	2.883	25032	1610	3.281	25032	161C	3.329
25032	160C	3.543	25032	161N	3.634	25032	161CB	3.892
25032	162N	4.376	25032	160CA	4.974		2750H2	4.982
25C33	1610	3.671	25C33	660	4.037	25C33	161C	4.309
25C33	66N	4.440	25C33	275OH2	4.717	25C33	161CA	4.785
25C33	65CA	4.801	25C34	660	2.806	25C34	66C	3.890
25C34	66N	4.077	25C34	26CB	4.273	25C34	66CA	4.590
25C34	1610	4.797	25C34	67N	4.862	25C34	26CG	4.869
25C34	163CB	4.955	25C35	660	3.565	25C35	209CD2	4.310
25C35	134CB	4.451	25C35	163CB	4.577	25C35	66C	4.668
25C35	67CD1	4.755	25C35		4.786	25C35	68SD	4.819
	1600					25C35		4.979
25C36	161C	3.690	25C36	1610				
	134CB						163N	3.971
	161CA							
25C36	161N	4.238						
	163CB			1620				
	160CB							
25C37	209CD2	3.499	25C37	67CD1	3.574	25C37	67CE1	3.696

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#### TABLE XXI

25C37 660	3.889	25C37	67CG	4.551	25C37	1600	4.714
25C37 67CZ	4.732	25C37	66C	4.737	25C37	134CB	4.814
25C37 67CA	4.821	25C37 2	09CG	4.994	25C38	65CA	3.869
25C38 1610	3.955	25C38	66N	4.001	25C38	660	4.219
25C38 26CD1	4.264	25C38	25SG	4.271	25C38	2750H2	4.287
25C38 65C	4.487	25C38	230	4.627	25C38	26CB	4.655
25C38 26CG	4.792	25C38	640	4.873	25C38	161C	4.899
25C38 65N	4.993	25039	65CA	2.703	25039	66N	3.332
25039 65C	3.531	25039 2	750H2	3.677	25039	640	3.733
25039 65N	3.795	25039	26CD1	4.043	25039	230	4.088
25039 64C	4.155	25039	660	4.358	25039	66CA	4.647
25039 650	4.732	25039	26CG	4.801	25039	26NE1	4.806
25039 1610	4.910	25039	66C	4.976	25N40	25SG	2.943
25N40 1610	3.660	25N40	26CD1	4.126	25 <b>N4</b> 0	26N	4.248
25N40 230	4.287	25N40	26CB	4.350	25N40	163N	4.551
25N40 25CB	4.560	25N40	26CG	4.646	25N40	161C	4.657
25N40 25N	4.685	25N40	65CA	4.705	25 <b>N4</b> 0	660	4.809
25N40 162CA	4.847	25N40	26CA	4.856	25 <b>N4</b> 0	6 <b>6N</b>	4.876
25N40 23C	4.933	25N40 1	.63CB	4.949	25 <b>N4</b> 0	25CA	4.979
25N40 25C	4.984	25C41	25SG	2.569	25C41	230	3.094
25C41 25N	3.498	25C41	23C	3.558	25C41	26CD1	3.732
25C41 26N	3.810	25C41	25CB	3.828	25C41	25CA	4.081
25C41 23CA	4.144	25C41	24N	4.195	25C41	24C	4.371
25C41 25C	4.413	25C41	26CB	4.509	25C41	26CG	4.528
25C41 24CA	4.533	25C41 1	610	4.584	25C41	65CA	4.716
25C41 26NE1	4.762	25C41	26CA	4.764	25N42	2750H2	3.989
25N42 1610	4.257	25N42	66N	4.573	25N42	660	4.641
25N42 65CA	4.726	25N42 1	161C	4.818	25N42	1600	4.862
25N42 161CA	4.871						

#### TABLE XXII

Table of distances in Ångstroms between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	184CB	4.217	25C1	1840	4.458	25C1	184CG	4.534
25C1	184CD2	4.719	25C1	184CE3	4.742	25C1	188CD1	4.982
25C2	1840	3.589	25C2	184CB	3.931	25C2	180D1	4.344
25C2	184C	4.365	25C2	184CA	4.384	25C2	184CG	4.474
25C3	180D1	3.467	25C3	184CB	3.731	25C3	1840	3.805
25C3	184CA	3.925	25C3	184CG	4.004	25C3	184C	4.224
25C3	184CD1	4.243	25C3	18CG	4.424	25C3	18ND2	4.653
25C3	184CD2	4.783	25C3	200	4.954	25C4	184CG	3.557
25C4	184CD1	3.557	25C4	184CB	3.839	25C4	184NE1	4.069
25C4	184CD2	4.096	25C4	184CA	4.267	25C4	180D1	4.278
25C4	184CE2	4.376	25C4	200	4.406	25C4	1840	4.793
25C4	184CE3	4.895	25C4	184C	4.990	25C5	184CG	3.622
25C5	184CD2	3.653	25C5	184CD1	3.870	25C5	184CE2	3.925
25C5	184NE1	4.046	25C5	184CB	4.116	25C5	184CE3	4.153
25C5	184CZ2	4.632	25C5	184CZ3	4.798	25C5	184CA	4.969
25C6	184CD2	4.011	25C6	184CE3	4.065	25C6	184CG	4.134
25C6	184CB	4.303	25C6	184CE2	4.616	25C6	184CZ3	4.700
25C6	184CD1	4.765	25C7	200	3.159	25C7	184CD1	3.353
25C7	19CG	3.702	25C7	184NE1	3.716	25C7	184CG	3.910
25 <b>C7</b>	20C	4.053	25C7	20 <b>N</b>	4.098	25C7	180D1	4.206
25C7	19CD	4.236	25 <b>C</b> 7	20CA	4.370	25C7	184CE2	4.426
25C7	184CB	4.474	25C7	19NE2	4.497	25C7	184CD2	4.546
25C7	184CA	4.592	25C7	1330	4.805	25C7	190E1	4.886
25C7	19C	4.994	2508	200	2.990	2508	20C	4.056
2508	184NE1	4.178	2508	184CD1	4.264	2508	19CG	4.312
2508	19NE2	4.344	2508	19CD	4.511	2508	184CE2	4.748
2508	20CA	4.839	2508	20N	4.842	2508	184CG	4.873
2508	21N	4.938	2508	21CA	4.989	25 <b>C</b> 9	200	3.137
25C9	19NE2	3.361	2509	19CD	3.886	25C9	19CG	4.087

#### TABLE XXII

25C9	184NE1	4.190	25C9	20C	4.350	25C9	220	4.421
25C9	184CD1	4.624	25C9	190E1	4.690	25C9	22N	4.828
25C9	184CE2	4.870	25C9	21CA	4.905	25C9	210E1	4.986
25010	200	2.532	25010	19NE2	2.700	25010	220	3.220
25010	19CD	3.446	25010	19CG	3.596	25010	22N	3.702
25010	20C	3.735	25010	22C	3.951	25010	21CA	4.210
25010	21C	4.416	25010	21N	4.441	25010	190E1	4.463
25010	22CA	4.519	25010	19CB	4.714	25010	20N	4.734
25010	23N	4.741	25010	20CA	4.838	25010	184NE1	4.848
25010	210E1	4.906	25010	23CA	4.996	25C11	19NE2	3.727
25C11	19CD	4.636	25C11	184NE1	4.779	25C11	220	4.911
25C12	19NE2	3.752	25C12	220	4.028	25C12	23CA	4.150
25C12	22C	4.246	25C12	23N	4.275	25C12	224OH2	4.684
25C12	22N	4.779	25C12	19CD	4.925	25C12	200	4.975
25C13	210E1	4.156	25C13	22N	4.627	25C13	22C	4.667
25C13	23N	4.712	25C13	220	4.742	25C13	21C	4.914
25C13	23CA	4.953	25C14	22N	3.473	25C14	21C	3.659
25C14	22C	3.759	25C14	210E1	3.793	25C14	23N	3.863
25C14	22CA	3.868	25C14	21CA	4.095	25C14	210	4.118
25C14	220	4.147	25C14	23CA	4.513	25C14	200	4.765
25C14	21CB	4.851	25C14	21CD	4.885	25C15	210E1	3.260
25C15	21CD	4.221	25C15	21NE2	4.650	25C15	21CA	4.882
25C15	200	4.924	25C16	19NE2	3.267	25C16	19CD	4.074
25C16	162ND1	4.145	25C16	190E1	4.238	25C16	162CE1	4.291
25C16	184NE1	4.295	25C16	184CZ2	4.540	25C16	184CE2	4.778
25C16	25SG	4.792	25C16	23CA	4.932	25C16	220	4.976
25017	19NE2	3.075	25017	184NE1	3.183	25017	162CE1	3.232
25017	162ND1	3.349	25017	190E1	3.394	25017	19CD	3.490
25017	184CZ2	3.632	25017	184CE2	3.732	25017	162NE2	4.167
25017	162CG	4.366	25017	184CD1	4.379	25017	25CB	4.553
25017	25 SG	4.612	25017	19CG	4.742	25017	162CD2	4.779
25017	184CH2	4.923	25N18	19NE2	3.859	25N18	25 <i>S</i> G	4.056
25N18	162ND1	4.100	25N18	1610	4.319	25N18	23CA	4.407
	2240H2		25N18	162CE1	4.601	25N18	19CD	4.788
25N18	25CB	4.834	25N18	190E1	4.862	25N19	25SG	2.787
25N19	162ND1	3.174	25N19	163.0	3.363	25N19	25CB	3.873
25N19	162CE1	3.953	25N19	162CA	4.146	25N19	162CG	4.165

# TABLE XXII

25N19	19NE2	4.334	25N19	162CB	4.414	25N19	161C	4.448
25N19	23CA	4.774	25N19	162N	4.806	25N19	190E1	4.807
25N19	224OH2	4.923	25N20	19NE2	3.800	25N20	184NE1	4.071
25N20	19CD	4.420	25 <b>N</b> 20	200	4.457	25N20	184CE2	4.475
25N20	184CZ2	4.573	25N20	184CD1	4.849	25N20	190E1	4.977
25N20	19CG	4.982	25C21	25SG	1.799	25C21	25CB	3.030
25C21	25N	3.707	25C21	162ND1	3.784	25C21	23CA	3.850
25C21	23C	3.907	25C21	230	3.946	25C21	25CA	3.958
25C21	19NE2	4.007	25C21	1610	4.284	25C21	162CE1	4.379
25C21	2240H2	4.436	25C21	24N	4.507	25C21	190E1	4.543
25C21	162CA	4.728	25C21	19CD	4.730	25C21	26N	4.758
25C21	25C	4.830	25C21	24C	4.870	25C21	162CG	4.947
25C21	163N	4.989	25022	25SG	2.443	25022	19NE2	2.871
25022	25CB	2.954	25022	23CA	3.128	25022	25N	3.144
25022	23C .	3.258	25022	25CA	3.641	25022	24N	3.649
25022	230	3.656	25022	19CD	3.696	25022	190E1	3.727
25022	220	4.038	25022	162ND1	4.175	25022	24C	4.265
25022	23N	4.327	25022	162CE1	4.422	25022	224OH2	4.423
25022	24CA	4.529	25022	22C	4.622	25022	25C	4.800
25022	26N	4.898	25C23	610D1	3.673	25C23	590	3.735
25C23	67CE2	4.205	25C23	67CD2	4.463	25C23	2640H2	4.473
25C23	61CG	4.499	25C23	610D2	4.706	25C23	59C	4.887
25C24	590	3.179	25C24	610D1	3.691	25C24	67CD2	3.717
25C24	60ND2	3.736	25C24	67CE2	3.832	25C24	60CA	3.942
25C24	59C	4.226	25C24	61CG	4.362	25C24	66CA	4.399
25C24	60C	4.521	25C24	61N	4.525	25C24	60N	4.567
25C24	700D1	4.595	25C24	610D2	4.702	25C24	60CG	4.710
25C24	60CB	4.837	25C24	67N	4.915	25C24	66C	4.929
25C24	650	4.975	25C24	67CG	4.986	25C25	66CA	3.301
25C25	610D1	3.379	25C25	67CD2	3.477	25C25	67CE2	3.538
25C25	60ND2	3.834	25C25	66N	3.859	25C25	650	3.860
25C25	66C	3.977	25C25	65C	4.082	25C25	60CA	4.110
25C25	61CG	4.139	25C25	61N	4.153	25C25	590	4.235
25C25	67N	4.302	25C25	60C	4.526	25C25	60CG	4.585
25C25	660	4.675	25C25	60CB	4 701	25C25	61CB	4.755
25C25	67CG	4.785	25C25	610D2	4.825	25C25	67CZ	4.879
25C26	610D1	3.031	25C26	67CE2	3.662	25C26	66CA	3.882

25C26	66N	3.954	25C26	650	4.020	25C26	65C	4.027
25C26	67CD2	4.058	25C26	61CG	4.066	25C26	66C	4.566
25C26	61CB	4.672	25C26	61N	4.764	25C26	67CZ	4.811
25C26	65CA	4.818	25C26	610D2	4.956	25C26	660	4.959
25C27	610D1	3.004	25C27	67CE2	4.045	25C27	61CG	4.208
25C27	67CD2	4.744	25C27	610D2	4.951	25C27	670H	4.988
25C28	610D1	3.343	25C28	67CE2	4.306	25C28	61CG	4.423
25C28	2640H2	4.506	25C28	610D2	4.829	25C28	67CD2	4.927
25C29	6 <b>6N</b>	3.077	25C29	65C	3.103	25C29	650	3.481
25C29	66CA	3.531	25C29	65CA	3.578	25C29	610D1	3.662
25C29	67CE2	4.189	25C29	640	4.192	25C29	66C	4.265
25C29	660	4.382	25C29	65N	4.522	25C29	67CD2	4.589
25C29	61CG	4.593	25C29	64C	4.738	25C29	61CB	4.762
25030	66N	3.458	25030	67CE2	3.803	25030	65C	3.840
25030	660	4.008	25030	66CA	4.009	25030	65CA	4.053
25030	66C	4.287	25030	67CD2	4.421	25030	67CZ	4.431
25030	670H	4.470	25030	650	4.551	25030	640	4.750
25C31	66N	3.498	25C31	65CA	3.654	25C31	65C	3.885
25C31	660	4.018	25C31	66CA	4.416	25C31	640	4.463
25C31	66C	4.608	25C31	650	4.872	25C31	67CE2	4.874
25C31	65N	4.883	25032	65CA	3.690	25032	640	3.833
25032	66N	4.273	25032	65C	4.343	25032	64C	4.667
25032	65N	4.681	25C33	660	3.876	25C33	66N	4.173
25C33	1610	4.383	25C33	65CA	4.417	25C33	161C	4.670
25C33	25 <b>SG</b>	4.802	25C33	65C	4.843	25C33	66C	4.902
25C34	660	3.833	25C34	161C	4.228	25C34	1610	4.287
25C34	162N	4.305	25C34	163N	4.452	25C34	162CA	4.613
25C34	162C	4.644	25C34	25SG	4.656	25C34	161CA	4.669
25C34	163CB	4.789	25C34	66N	4.994	25C35	660	2.981
25C35	66C	4.208	25C35	163CB	4.767	25C35	26CB	4.871
25C35	66N	4.887	25C35	67CA	4.934	25C35	209CD2	4.958
25C35	68SD	4.962	25C35	67CD1	4.992	25C35	67CE1	5.000
25C36	660	3.430	25C36	68SD	3.532	25C36	163CB	3.585
25C36	163CA	4.250	25C36	134CB	4.323	25C36	163N	4.364
25C36	26CB	4.411	25C36	68CE	4.444	25C36	209CD2	4.518
25C36	66C	4.590	25C36	57CA	4.718	25C36	152C	4.845
25C36	26CX	4.975	25C37	660	3.900	25C37	67CE1	4.120

								4 430
25C37 2	09CD2	4.133	25C37	67CZ	4.304	25C37	67CD1	4.472
25C37	670H	4.559	25C37	1600	4.722	25C37	67CE2	4.812
	34CB	4.928	25C37	67CG	4.960	25C38	65CA	3.685
25C38	25SG	3.703	25C38	66N	3.862	25C38	660	4.169
25C38	26CD1	4.241	25C38	65C	4.328	25C38	230	4.331
	1610	4.500	25C38	2240H2	4.706	25C38	26CB	4.746
25C38	65N	4.855	25C38	26CG	4.876	25039	66N	2.954
25039	26CD1	3.044	25039	65CA	3.072	25039	65C	3.468
25039	660	3.555	25039	230	3.635	25039	26CG	3.715
25039	26CB	3.796	25039	25 <b>S</b> G	3.860	25039	26NE1	4.093
25039	66CA	4.102	25039	66C	4.239	25039	65N	4.292
25039	26N	4.359	25039	650	4.668	25039	23C	4.669
25039	26CX	4.689	25039	2240H2	4.850	25039	26CD2	4.970
25039 25N40	25SG	3.113	25N40	1610	3.678	25N40	224OH2	4.045
	230	4.202	25N40	65CA	4.334	25N40	161C	4.525
25N40	23C	4.779	25N40	23CA	4.821	25N40	162CA	4.878
25N40	25CB	4.902	25N40	26CD1	4.925	25N41	25SG	2.621
25N40	230	3.122	25N41	2240H2	3.404	25N41	23C	3.447
25N41	23CA	3.458	25N41	25CB	4.118	25N41	25N	4.161
25N41		4.403	25N41	65CA	4.453	25N41	1610	4.531
25N41	24N	4.572	25N41		4.709	25 <b>N4</b> 1	26N	4.861
25N41	26CD1	4.898	25N42		3.140	25N42	66N	3.340
25N41	23N		25N42		4.005	25 <b>N4</b> 2	65C	4.084
25N42	65CA	3.978 4.168	25N42		4.883			
2 E NT # 2	6652	4 100	4 J I Y Y 4	, 0,022				

#### **TABLE XXIII**

Table of distances in Angstroms between atoms of the inhibitor and protein for all protein atoms within 5 Angstroms of the inhibitor (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	184CB	4.236	25C1	184CG	4.418	25C1	1840	4.425
25C1	184CD2	4.593	25C1	184CE3	4.660	25C1	188CD1	4.721
25C2	1840	3.268	25C2	184CB	3.689	25C2	184C	4.080
25C2	184CG	4.182	25C2	184CA	4.182	25C2	184CD2	4.784
25C2	184CD1	4.821	25C2	18ND2	4.999	25C3	1840	3.481
25C3	184CB	3.729	25C3	184CA	3.863	25C3	184C	3.967
25C3	180D1	4.001	25C3	184CG	4.006	25C3	184CD1	4.249
25C3	18ND2	4.280	25 <b>C</b> 3	18CG	4.516	25C3	184CD2	4.834
25C4	184CD1	4.005	25C4	184CG	4.080	25C4	184CB	4.302
25C4	180D1	4.440	25C4	200	4.528	25C4	184CA	4.558
25C4	184NE1	4.584	25C4	184CD2	4.705	25C4	1840	4.729
25C4	184CE2	4.985	25C5	184CG	4.312	25C5	184CD1	4.376
25C5	184CD2	4.504	25C5	184NE1	4.612	25 <b>C</b> 5	184CE2	4.694
25C5	184CB	4.766	25C5	21NE2	4.820	25C6	184CD2	4.450
25C6	184CG	4.480	25C6	184CE3	4.643	25C6	184CB	4.744
25C6	1430E1	4.804	25C6	184CE2	4.925	25C6	184CD1	4.935
25C7	200	3.040	25C7	20C	3.594	25C7	20CA	3.795
25C7	20N	3.853	25C7	180D1	3.864	25C7	184CD1	4.135
25 <b>C</b> 7	19CG	4.463	25C7	21NE2	4.595	25C7	21N	4.617
25C7	184NE1	4.641	25C7	184CG	4.659	25C7	18CG	4.887
25C7	19C	4.937	25C7	184CA	4.958	2508	200	2.667
2508	19CG	3.422	2508	134CD1	3.450	2508	20C	3.631
2508	184NE1	3.675	2508	20N	3.871	2508	19CD	3.993
2508	20CA	4.102	2508	190E1	4.232	2508	184CG	4.325
2508	180D1	4.411	2508	184CE2	4.622	2508	19C	4.642
2508	21N	4.725	2508	19CB	4.727	2508	19NE2	4.771
2508	1830	4.826	2508	184CD2	4.974	25C9	200	3.188
25C9	184NE1	3.536	25C9	184CD1	3.844	25C9	19CG	4.035

25C9	19CD	4.106	25C9	190E1	4.229	25C9	20C	4.321
25C9	184CE2	4.325	25C9	19NE2	4.650	25C9	184CG	4.743
25C9	184CZ2	4.957	25C9	21NE2	4.999	25010	184NE1	3.675
25010	184CE2	4.068	25010	184CD1	4.137	25010	200	4.244
25010	184CZ2	4.481	25010	184CD2	4.713	25010	184CG	4.754
25010	21NE2	4.824	25C11	19NE2	4.159	25C11	220	4.229
25C11	19CD	4.273	25C11	200	4.296	25C11	190E1	4.447
25C11	184NE1	4.589	25C11	22C	4.735	25C11	23CA	4.867
25C11	19CG	4.868	25C11	22N	4.974	25C11	23N	4.979
25C12	220	3.890	25C12	22C	3.995	25C12	22N	4.110
25C12	23N	4.111	25C12	23CA	4.306	25C12	200	4.346
25C12	21C	4.588	25C12	22CA	4.594	25C12	210E1	4.629
25C12	21CA	4.715	25C12	19NE2	4.762	25C13	210E1	3.498
25C13	21CD	4.390	25C13	21CA	4.587	25C13	22N	4.615
25C13	21C	4.659	25C13	200	4.898	25C13	21NE2	4.928
25C14	210E1	2.922	25C14	21C	3.654	25C14	21CA	3.785
25C14	22N	3.919	25C14	21CD	3.931	25C14	210	3.962
25C14	21CB	4.117	25C14	22CA	4.561	25C14	21CG	4.659
25C14	22C	4.749	25C14	21NE2	4.841	25C14	23N	4.886
25C14	200	4.941	25C15	210E1	2.917	25C15	21CD	3.612
25C15	21NE2	3.794	25C15	21CA	4.698	25C15	200	4.791
25C15	21CG	4.892	25C16	19NE2	3.649	25C16	190E1	3.851
25C16	19CD	3.913	25C16	23CA	4.390	25C16	184NE1	4.396
25C16	220	4.401	25C16	162ND1	4.744	25C16	19CG	4.960
25C16	23N	4.963	25C16	22C	4.981	25C16	184CZ2	4.990
25\$17	162ND1	3.565	25S17	184CZ2	3.585	25S17	184NE1	3.699
25 <b>S</b> 17	1.62CE1	3.945	25 <b>S</b> 17	190El	3.950	25517	184CE2	3.994
25 <b>S</b> 17	162CG	4.327	25517	19CD	4.502	25517	19NE2	4.573
25 <b>S</b> 17	162CB	4.756	25517	184CH2	4.771	25 <b>S</b> 17	162NE2	4.811
25\$17	184CD1	4.996	25N18	19NE2	3.269	25N18	23CA	3.310
25N18	19CD	3.941	25N18	220	3.976	25N18	190E1	4.019
25N18	23C	4.156	25N18	25 <b>5</b> G	4.177	25N18	23N	4.187
25N18	22C	4.455	25N18	25CB	4.460	25N18	24N	4.557
25N18	152ND1	4.710	25N18	25N	4.758	25N18	230	4.840
25C19	25SG	3.024	25C19	25CB	3.565	25C19	19NE2	3.784
25C19	162ND1	3.801	25C19	23CA	3.948	25C19	190E1	4.215
25C19	1610	4.243	25C19	19CD	4.411	25C19	25N	4.416

25C19	23C	4.434	25C19	162CE1	4.494	25C19	25CA	4.626
25C19	230	4.822	25C19	162CG	4.842	25C19	24N	4.896
25C19	162CA	4.932	25N20	200	3.090	25N20	19CD	3.488
25N20	19NE2	3.690	25N20	19CG	3.707	25N20	190E1	3.805
25N20	220	3.918	25N20	184NE1	3.976	25N20	20C	4.316
25N20	184CD1	4.488	25N20	22N	4.551	25N20	22C	4.673
25N20	21CA	4.853	25N20	184CE2	4.906	25N20	19CB	4.997
25C21	162ND1	2.861	25C21	25SG	3.627	25C21	162CE1	3.661
25C21	162CG	3.744	25C21	1610	3.811	25C21	25CB	3.895
25C21	162CB	3.972	25C21	190E1	4.135	25C21	162CA	4.169
25C21	19NE2	4.408	25C21	19CD	4.664	25C21	162NE2	4.724
25C21	161C	4.759	25C21	184CZ2	4.788	25C21	162CD2	4.789
25C21	184NE1	4.807	25C21	162N	4.963	25C22	25SG	1.806
25C22	25CB	3.017	25C22	25N	3.658	25C22	23CA	3.756
25C22	23C	3.777	25C22	230	3.783	25C22	25CA	3.942
25C22	1610	4.202	25C22	19NE2	4.263	25C22	24N	4.389
25C22	162ND1	4.411	25C22	26N	4.767	25C22	24C	4.797
25C22	25C	4.896	25C22	162CA	4.914	25C22	26CD1	4.985
25023	25 <b>S</b> G	2.263	25023	23C	2.724	25023	25N	2.817
25023	23CA	2.863	25023	230	2.953	25023	25CB	2.966
25023	24N	3.216	25023	19NE2	3.371	25023	25CA	3.468
25023	24C	3.844	25023	24CA	4.001	25023	23N	4.229
25023	220	4.346	25023	19CD	4.418	25023	26N	4.527
25023	25C	4.570	25023	190E1	4.594	25023	26CD1	4.647
25023	22C	4.750	25023	162ND1	4.926	25023	240	4.984
25C24	640	4.232	25C24	610D1	4.886	25C25	640	2.914
25C25	610D1	4.012	25C25	64C	4.121	25C25	65CA	4.508
25C25	61CG	4.803	25C25	65N	4.812	25C26	640	3.004
25C26	610D1	3.284	25C26	65CA	3.778	25C26	65C	4.006
25C26	64C	4.052	25C26	66N	4.292	25C26	61CG	4.320
25C26	65N	4.406	25C26	650	4.498	25C26	61CB	4.929
25C27	610D1	3.654	25C27	640	4.372	25C27	66N	4.515
25C27	67CE2	4.579	25C27	65C	4.592	25C27	65CA	4.693
25C27	61CG	4.839	25C27	670H	4.900	25028	610D1	4.602
25C28	670H	4.61.4	25C28	67CE2	4.849	25C30	67CE2	3.326
25C30	66N	3.711	25C30	610D1	3.817	25C30	66CA	3.888
25C30	67CD2	3.908	25C30	67CZ	4.039	25C30	65C	4.143

25C30	670H	4.216	25C30	66C	4.303	25C30	660	4.459
25C30	650	4.648	25C30	65CA	4.662	25C30	67CG	4.980
25031	67CE2	2.958	25031	67CZ	3.233	25031	670H	3.437
25031	67CD2	3.606	25031	660	3.722	25031	66N	3.758
25031	66C	3.942	25031	66CA	3.994	25031	67CE1	4.053
25031	67CG	4.365	25031	65C	4.521	25031	67CD1	4.543
25031	67N	4.754	25031	65CA	4.909	25C32	660	3.869
25C32	66N	3.875	25C32	67CZ	4.120	25C32	670H	4.127
25C32	67CE2	4.188	25C32	66C	4.431	25C32	66CA	4.512
25C32	65CA	4.559	25C32	65C	4.577	25C32	67CE1	4.683
25C32	67CD2	4.797	25033	670H	4.349	25033	67CZ	4.706
25033	66N	4.896	25033	67CE2	4.948	25033	1600	4.987
25C34	660	3.926	25C34	6 <b>6N</b>	4.154	25C34	1610	4.291
25C34	65CA	4.393	25C34	161C	4.680	25C34	25 <b>S</b> G	4.728
25C34	1600	4.762	25C34	65C	4.835	25C34	66C	4.889
25C35	1610	3.923	25C35	1600	3.934	25C35	161C	3.950
25C35	660	4.195	25C35	162N	4.296	25C35	161CA	4.330
25C35	163N	4.545	25C35	160C	4.652	25C35	25 <b>SG</b>	4.661
25C35	162CA	4.717	25C35	161N	4.821	25C35	162C	4.844
25C36	660	3.516	25C36	163CB	3.928	25C36	163N	4.123
25C36	163CA	4.451	25C36	134CB	4.614	25C36	162C	4.695
25C36	6 <b>6</b> C	4.739	25C36	25SG	4.777	25C36	1600	4.790
25C36	68SD	4.793	25C36	162N	4.890	25C36	161C	4.930
25C36	26CB	4.984	25C37	660	3.229	25C37	67CD1	3.782
25C37	67CE1	3.814	25C37	209CD2	4.109	25C37	66C	4.351
25C37	134CB	4.449	25C37	67CG	4.461	25 <b>C</b> 37	67CZ	4.520
25C37	67CA	4.594	25C37	68SD	4.609	25C37	163CB	4.822
25C37	68CE	4.913	25C37	67N	4.944	25C37	1600	4.999
25C38	6 <b>6</b> 0	2.883	25C38	163CB	3.258	25 <b>C</b> 38	26CB	3.517
25C38	25SG	3.915	25C38	163N	4.010	25 <b>C</b> 38	66C	4.045
25C38	26CA	4.069	25C38	25N	4.161	25C38	163CA	4.178
25C38	68SD	4.184	25C38	26CG	4.396	25C38	26CD1	4.512
25C38	6 <b>6N</b>	4.558	25C38	67CA	4.916	25C38	67N	4.916
25C38	66CA	4.920	25C38	162C	4.937	25C39	65CA	3.532
25C39	25 <b>S</b> G	3.557	25C39	66N	3.788	25C39	660	4.161
25C39	1610	4.207	25C39	65C	4.218	25C39		4.481
25C39	65N	4.564	25C39	161C	4.925	25C39	230	4.951

25C39	66CA	4.972	25040	66N	2.864	25040	65CA	2.895
25040	26CD1	3.289	25040	65C	3.326	25040	25 <i>S</i> G	3.487
25040	660	3.497	25040	66CA	4.034	25040	26CG	4.074
25040	65N	4.098	25040	26NE1	4.177	25040	66C	4.186
25040	230	4.205	25040	26CB	4.291	25040	650	4.516
25040	26N	4.644	25 <b>N4</b> 1	25 <i>S</i> G	3.236	25N41	1610	3.370
25N41	65CA	4.079	25N41	161C	4.343	25N41	230	4.777
25N41	65N	4.879	25 <b>N4</b> 1	66N	4.926	25N42	25SG	2.820
25N42	230	3.597	25 <b>N4</b> 2	23CA	3.792	25N42	23C	3.925
25N42	65CA	4.086	25N42	1610	4.156	25N42	25CB	4.397
25N42	65N	4.514	25N42	25 <b>N</b>	4.705	25N42	26CD1	4.802
25N42	24N	4.905	25N43	660	3.027	25N43	66N	3.312
25N43	66C	3.828	25N43	66CA	4.057	25N43	65CA	4.128
25N43	65C	4.171	25 <b>N4</b> 3	67CZ	4.741	25N43	67CE2	4.807
25N43	67CF1	4 973	25N43	67N	4 980			

# TABLE XXIV

Table of distances in Angstroms between atoms of the inhibitor and protein for all protein atoms within 5 Angstroms of the inhibitor 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	2410H2	3.570	25C1	184CD1	3.752	25C1	184CB	3.771
25C1	184CG	3.795	25C1	1840	3.855	25C1	180D1	3.918
25C1	184CA	3.940	25C1	18ND2	4.077	25C1	184C	4.295
25C1	18CG	4.335	25C1	200	4.526	25C1	184NE1	4.545
25C1	184CD2	4.638	25C2	180D1	2.991	25C2	184CD1	3.364
25C2	184CA	3.673	25C2	200	3.682	25C2	18CG	3.691
25C2	184CG	3.828	25C2	20N	3.856	25C2	18ND2	3.872
25C2	184CB	3.927	25C2	1840	4.146	25C2	19CG	4.202
25C2	184C	4.236	25C2	184NE1	4.236	25C2	1830	4.241
25C2	20CA	4.325	25C2	20C	4.444	25C2	19N	4.506
25C2	18 <b>4N</b>	4.779	25C2	2410H2	4.781	25C2	19C	4.859
25C2	184CD2	4.882	25C2	183C	4.942	25C2	18CB	4.966
25C3	200	2.766	25C3	184CD1	3.449	25C3	19CG	3.485
25C3	20N	3.566	25C3	180D1	3.700	25C3	20C	3.715
25C3	184NE1	3.945	25C3	20CA	3.980	25C3	19CD	4.274
25C3	184CG	4.298	25C3	19C	4.413	25C3	1830	4.573
25C3	19N	4.586	25C3	184CA	4.627	25C3	19CB	4.636
25C3	18CG	4.663	25C3	19CA	4.734	25C3	19NE2	4.739
25C3	184CB	4.805	25C3	190E1	4.890	25C3	21N	4.941
25C3	184CE2	4.965	25C4	200	2.926	25C4	184CD1	3.912
25C4	184NE1	3.991	25C4	20C	4.103	25C4	19CG	4.382
25C4	20N	4.662	25C4	184CG	4.708	25C4	20CA	4.787
25C4	19CD	4.799	25C4	184CE2	4.822	25C4	180D1	4.987
25C4	2410H2	4.989	25C4	19NE2	4.989	25C5	2410H2	3.832
25C5	200	3.929	25C5	184CD1	4.241	25C5	184NE1	4.312
25C5	184CG	4.671	25C5	184CE2	4.778	25C5	184CD2	4.987
25C6	2410H2	2.940	25C6	194CD1	4.176	25C6	184CG	4.243
25C6	184CB	4.555	25C6	184NEL	4.583	25C6	200	4.631
25C6	184CD2	4.694	25C6	184CE2	4.889	25C7	200	2.932

## **TABLE XXIV**

25C7	20C	4.097	25C7	19NE2	4.319	25C7	19CG	4.452
25C7	184NE1	4.478	25C7	19CD	4.533	25C7	21CA	4.627
25C7	220	4.737	25C7	184CD1	4.756	25C7	21N	4.840
25C7	21C	4.925	2508	19NE2	3.416	2508	19CD	3.782
2508	184NE1	3.951	2508	200	4.046	2508	19CG	4.180
2508	190E1	4.351	2508	220	4.445	2508	184CD1	4.559
2508	184CE2	4.877	25C9	19NE2	4.211	25C9	184NE1	4.403
25C9	19CD	4.724	25C11	162ND1	4.502	25C11	184CZ2	4.687
25C11	162CE1	4.868	25C14	162ND1	3.357	25C14	162CG	3.831
25C14	162CB	4.001	25C14	162CE1	4.036	25C14	162CA	4.368
25C14	1610	4.489	25C14	184CZ2	4.614	25C14	162CD2	4.682
25C14	1610D1	4.720	25C14	162N	4.758	25C14	162NE2	4.766
25C14	161C	4.794	25C14	25SG	4.940	25015	162CB	3.130
25015	162ND1	3.165	25015	162CG	3.174	25015	1610D1	3.790
25015	162CA	3.794	25015	162CE1	3.969	25015	162CD2	4.004
25015	162N	4.125	25015	184CZ2	4.191	25015	162NE2	4.407
25015	161C	4.425	25015	1610	4.436	25015	137CB	4.597
25015	161CG	4.704	25015	1370	4.758	25015	161CB	4.862
25015	184CH2	4.953	25N16	162ND1	3.293	25N16	25SG	3.828
25N16	1610	3.898	25 <b>N</b> 16	162CE1	4.030	25N16	162CG	4.119
25N16	162CA	4.252	25N16	162CB	4.323	25N16	161C	4.537
25N16	25CB	4.583	25N16	162N	4.723	25N16	19NE2	4.776
25N17	25 <b>S</b> G	2.713	25N17	1610	2.829	25N17	162ND1	2.962
25N17	162CA	3.321	25N17	161C	3.640	25N17	162CG	3.800
25N17	162CB	3.840	25N17	162N	3.876	25N17	25CB	3.878
25N17	162CE1	3.946	25N17	163N	4.346	25N17	162C	4.395
25N17	161CA	4.922	25N17	162CD2	4.988	25C18	184NE1	4.080
25C18	19NE2	4.103	25C18	184CZ2	4.414	25C18	162ND1	4.448
25C18	162CE1	4.452	25C18	19CD	4.549	25C18	190E1	4.597
25C18	184CE2	4.604	25C19	25SG	1.793	25C19	25CB	3.071
25C19	1610	3.652	25C19	162ND1	3.659	25C19	25N	4.072
25C19	25CA	4.137	25C19	230	4.303	25C19	162CE1	4.350
25C19	162CA	4.352	25C19	19NE2	4.435	25C19	23C	4.511
25C19	163N	4.621	25C19	23CA	4.646	25C19	161C	4.687
25C19	162CG	4.766	25C19	190E1	4.843	25C19	25C	4.879
25C19	26N	4.949	25020	25SG	2.529	25020	25CB	3.171
25020	19NE2	3.371	25020	23CA	3.588	25020	23C	3.681

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# TABLE XXIV

25020	25N	3.725	25020	230	3.770	25020	162ND1	4.071
25020	25CA	4.076	25020	190E1	4.231	25020	19CD	4.248
25020	24N	4.306	25020	162CE1	4.450	25020	1610	4.781
25020	24C	4.875	25020	23N	4.969	25C21	160CD1	3.590
25C21	1580	3.615	25C21	160CG	3.707	25C21	160CB	3.983
25C21	160N	4.442	25C21	158C	4.580	25C21	160CA	4.835
25C22	160CD1	3.593	25C22	160CG	4.157	25C22	160CB	4.282
25C22	209CD2	4.511	25C22	209CD1	4.980	25C22	1580	4.986
25C23	160CD1	4.157	25C23	209CD2	4.159	25C23	67CE1	4.205
25C23	160CB	4.279	25C23	1600	4.519	25C23	670H	4.588
25C23	160CG	4.625	25C23	67CZ	4.894	25C24	1600	3.474
25C24	160CB	3.993	25C24	160C	4.419	25C24	160CA	4.596
25C24	160N	4.617	25C24	160CD1	4.665	25C24	160CG	4.704
25C24	67CE1	4.808	25C24	670H	4.862	25C25	1600	3.271
25C25	160N	3.593	25C25	160CB	3.664	25C25	1580	3.952
25C25	160CA	3.958	25C25	160C	3.985	25C25	160CG	4.302
25C25	160CD1	4.653	25C25	159C	4.710	25C25	159CA	4.972
25C25	158C	4.995	25C26	1580	2.899	25C26	160N	3.477
25C26	160CB	3.661	25C26	160CG	3.791	25C26	158C	3.973
25C26	160CA	4.096	25C26	160CD1	4.159	25C26	1600	4.202
25C26	159C	4.497	25C26	159CA	4.565	25C26	160C	4.606
25C26	159N	4.714	25C26	158CA	4.955	25C27	1600	3.338
25C27	670H	4.363	25C27	67CE1	4.376	25C27	160C	4.512
25C27	67CZ	4.684	25C27	160CB	4.754	25028	1600	2.420
25028	160C	3.644	25028	161CA	4.394	25028	160CB	4.473
25028	161N	4.476	25028	16CCA	4.624	25028	1610	4.916
25028	161C	4.988	25C29	1600	3.266	25C29	160C	4.372
25C29	1610	4.438	25C29	161CA	4.632	25C29	161C	4.791
25C29	660	4.888	25C29	67CE1	4.893	25C29	161N	4.959
25030	67CE1	3.985	25030	660	4.035	25030	67CD1	4.256
25030	67CZ	4.313	25030	1600	4.446	25030	66C	4.682
25030	670H	4.687	25030	67CG	4.823	25030	67CE2	4.858
25C31	1610	3.206	25C31	161C	3.941	25C31	660	3.994
25C31	163CB	4.458	25C31	1.63N	4.486	25C31	25 <b>S</b> G	4.527
25C31	1600	4.531	25C31	1.61.CA	4.598	25C31	162N	4.697
25C31	162C	4.753	25C31	162CA	4.857	25C32	660	2.742
25C32	66C	3.901	25C32	163CB	4.205	25C32	26CB	4.317

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# TABLE XXIV

25C32	2 66N	4.512	25C32	67CA	4.651	25C32	1610	4.691
25C32	2 67N	4.724	25C32	68SD	4.771	25C32	66CA	4.830
25C33	660	3.447	25C33	68SD	3.722	25C33	163CB	3.798
25C33	3 209CD2	4.102	25C33	68CE	4.212	25C33	134CB	4.330
25C33	67CA	4.465	25C33	66C	4.538	25C33	67CD1	4.678
25C33	163CA	4.689	25C33	26CB	4.751	25C33	163N	4.950
25C34	134CB	3.099	25C34 1	.63CB	3.583	25C34	209CD2	3.879
25C34	134CA	3.973	25C34 1	.63CA	3.984	25C34	163N	4.154
25C34	68SD	4.317	25C34 1	620	4.344	25C34	162C	4.346
25C34	68CE	4.364	25C34 1	600	4.704	25C34	1610	4.819
25C34	660	4.910	25C34 1	61C	4.924	25C34	1330	4.972
25C34	134C	4.977	25C34 1	34N	4.991	25C35	209CD2	3.174
25C35	67CD1	3.240	25C35	660	3.588	25C35	67CA	3.813
25C35	67CE1	3.871	25C35	67CG	3.924	25C35	67CB	4.168
25C35	68SD	4.179	25C35 2	09CG	4.360	25C35	68CE	4.389
25C35	66C	4.393	25C35 2	340H2	4.424	25C35	67N	4.535
25C35	134CB	4.642	25C35	68N	4.661	25C35	67C	4.803
25C35	67CZ	4.930	25C35	67CD2	4.976	25C36	1610	3.232
25C36	25SG	3.829	25C36	660	4.065	25C36	161C	4.303
25C36	66N	4.338	25C36	65CA	4.529	25C36	26CD1	4.948
25C36	163N	4.963	25C36	26CB	5.000	25037	66N	3.256
25037	65CA	3.411	25037	660	3.590	25037	65C	3.845
25037	1610	4.314	25037	66CA	4.368	25037	66C	4.383
25037	26CD1	4.400	25037	25 <b>S</b> G	4.578	25037	65N	4.736
25037	640	4.804	25037	26CB	4.911	25037	26CG	4.987
25N38	1610	2.583	25N38	25 <b>S</b> G	2.797	25N38	161C	3.765
25N38	162CA	4.301	25N38 1	63N	4.348	25N38	162N	4.497
25N38	25CB	4.579	25N38 1	62C	4.720	25N38	161CA	4.814
25N38	163CB	4.921	25N39	25 <i>S</i> G	2.623	25N39	1610	3.421
25N39	230	3.947	25N39	25CB	4.194	25N39	23C	4.527
25N39	161C	4.625	25N39	85CA	4.670	25N39	25N	4.673
25N39	23CA	4.780	25N39 2	25CA	4.961	25N39	162CA	4.964
25N40	1600	3.177	25N40 16	510	3.216	25 <b>N4</b> 0	161C	3.641
25N40	161CA	3.804	25N40 16	50C	4.036	25 <b>N4</b> 0	161N	4.304
25N40	162N	4.578	25N40 6	560	4.918			

#### **TABLE XXV**

Table of distances in Angstroms between atoms of the inhibitor and protein for all protein atoms within 5 Angstroms of the inhibitor 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone.

Atom 1	Atom 2	Dist.	Atom 1	Atom 2	Dist.	Atom 1	Atom 2	Dist.
25C1 1	840						184CG	4.294
25C1 1	84CB	4.338	25C1	184CA		25C1	180D1	4.581
25C1 1	84C	4.634	25C1	184NE1	4.872	25C1	184CD2	4.992
25C2	200	3.689	25C2	20C	4.194	25C2	21NE2	4.254
25C2 1	84CD1	4.266	25C2	20N		25C2	20CA	4.394
25C2	19CG	4.696	25C2	180D1		25C2	184NE1	4.731
25C2 1	84CG	4.742	25C2	184CA	4.975	25C3	200	3.755
25C3 1	84CD1	4.039	25C3	184NE1	4.087	25C3	21NE2	4.111
25C3	20C	4.569	25C3	19CG	4.678	25C3	184CG	4.714
25C3 1	84CE2	4.790	25C3	19CD	4.992	25C4	184NE1	3.514
25C4 1	84CD1	3.761	25C4	184CE2	3.871	25C4	184CG	4.236
25C4 1	84CD2	4.302	25C4	184CZ2	4.446	25C4	21NE2	4.858
25C5 1	84CE2	3.679	25C5	184NE1	3.687	25C5	184CD2	3.688
25C5 1	84CD1	3.704	25C5	184CG	3.711	25C5	184CZ2	4.317
25C5 1	84CE3	4.332	25C5	184CB	4.417	25C5	184CH2	4.847
25C5 1	84CZ3	4.856	2 <b>5</b> C6	184CG	3.747	25C6	184CD1	3.955
25C6 1	84CB	3.964	25C6	184CD2	4.101	25C6	1840	4.262
25C6 1	84NE1	4.395	25C6	184CE2	4.487	25C6	184CA	4.531
25C6 1	84CE3	4.701	25C6	184C	4.882	25C7	184NE1	3.584
25C7 1	84CE2	3.808	25C7	184CZ2	3.945	25C7	184CD1	4.327
25C7 1	84CD2	4.651	25C7	184CH2	4.855	25C7	184CG	4.925
2508 1	84NE1	3.393	2508	184CE2	3.944	2508	184CZ2	4.069
2508 1	84CD1	4.312	2508	19NE2	4.602	2508	19CD	4.825
25C9 1	84NE1	3.291	25C9	184CZ2	3.627	25C9	184CE2	3.765
25C9 1	62ND1	3.949	25C9	162CE1	4.256	25C9	184CD1	4.478
25C9	19NE2	4.492	25C9	190E1	4.778	25C9	19CD	4.791
25C9 1	84CH2	4.870	25010	184NE1	2.688	25010	162ND1	2.878
25010 1	62CE1	3.073	25010	184CE2	3.353	25010	184CZ2	3.407
25010	190E1	3.804	25010	184CD1	3.882	25010	19NE2	3.911
25010	19CD	4.053	25010	162CG	4.083	25010	162NE2	4.247
25010 1	.84CD2	4.680	25010	184CH2	4.748	25010	162CD2	4.767
	62CB	4.822	25010	25CB	4.894	25010	184CG	4.921
25010	25SG	4.959	25C11	1610	3.603	25C11	162ND1	3.785

## TABLE XXV

25C11	162CB	4.170	25C11	162CG	4.352	25C11	184CZ2	4.547
25C11	161C	4.651	25C11	1610D1	4.678	25C11	162CE1	4.739
25C11	162CA	4.759	25C12	1610	3.598	25C12	1610D1	3.692
25C12	161CG	4.426	25C12	161C	4.497	25C12	162CB	4.500
25C12	161CB	4.644	25C12	162ND1	4.828	25C13	1610D1	3.668
25C13	1370	3.927	25C13	137C	4.080	25C13	138N	4.272
25C13	184CZ2	4.318	25C13	138CA	4.377	25C13	137CB	4.500
25C13	143NE2	4.527	25C13	161CG	4.603	25C13	1610	4.651
25C13	184CH2	4.656	25C13	162CB	4.732	25C13	137CA	4.757
25C14	143NE2	3.452	25C14	184CZ2	4.059	25C14	1370	4.191
25C14	184CH2	4.327	25C14	143CD	4.676	25C14	137C	4.726
25C14	138CA	4.757	25C14	138N	4.988	25C15	1610D1	3.025
25C15	138CA	3.182	25C15	138N	3.231	25C15	137C	3.477
25C15	138CB	3.598	25C15	1370	3.616	25C15	161CG	3.781
25C15	137CA	4.362	25C15	161ND2	4.367	25C15	137N	4.495
25C15	138C	4.523	25C15	137CB	4.599	25C15	143NE2	4.610
25C15	1380G	4.662	25C15	161CB	4.679	25C15	1380	4.905
25C15	1610	4.924	25C16	1610	3.101	25C16	162ND1	4.231
25C16	161C	4.310	25C16	162CB	4.562	25C16	25 <b>S</b> G	4.612
25C16	162CA	4.657	25C16	162CG	4.876	25C16	162N	4.992
25017	1610	3.437	25017	161C	4.661	25N18	1610	3.143
25N18	25SG	3.301	25N18	162ND1	3.608		162CA	4.088
25N18	162CB	4.239	25N18	161C	4.252		162CG	4.379
25N18	25CB	4.462	25N18	162CE1	4.653	25N18	162N	4.681
25N18	19NE2	4.948	25N18	23CA	4.995	25C19	25SG	2.850
25C19	1610	3.592	25C19	23CA	4.261	25C19	25CB	4.284
25C19	162ND1	4.530	25C19	23C	4.535	25C19	162CA	4.545
25C19	161C	4.613	25C19	230	4.725	25C19	25N	4.814
25N20	184CZ2	4.159	25N20		4.342		184NE1	4.452
25N20	184CE2	4.669	25N20	162CE1		25C21		3.074
25C21	25SG	3.220	25C21	161C	3.894		162CA	4.288
25C21	162N	4.439	25C21	65CA	4.747		161CA	4.860
25C21	25CB	4.980	25C22		1.762	25C22		2.996
25C22	25N	3.317	25C22	25CA	3.720	25C22	23C	3.769
25C22	23CA	3.914	25C22	230	4.043	25C22	24N	4.052
25C22	19NE2	4.196	25C22	24C	4.394	25C22		4.399
25C22	26N	4.510	25C22	25C	4.601	25C22		4.652
25C22	24CA	4.697		162CA	4.792		163N	4.837
25C22	26CD1	4.881	25023	25SG	2.430	25023	25N	2.866
25023	25CB	2.896	25023	19NE2	3.003	25023		3.259
25023	23C	3.263	25023	24N	3.328	25023		3.459
25023	230	3.894	25023	24C	3.945	25023		4.004
25023	190E1	4.163	25023	24CA	4.166	25023	220	4.360

## TABLE XXV

25023	162ND1	4.426	25023	23N	4.535	25023	25C	4.653
25023	26N	4.784	25023	22C	4.908	25023	162CE1	4.912
25C24	590	3.356	25C24	610D2	4.112	25C24	60CA	4.139
25C24	61N	4.252	25C24	60ND2	4.327	25C24	60C	4.397
25C24	67CE2	4.413	25C24	59C	4.424	25C24	67CD2	4.649
25C24	61CB	4.711	25C24	60N	4.781	25C24	650	4.810
25C24	66CA	4.888	25C24	61CG	4.890	25C25	610D2	2.979
25C25	61CB	3.456	25C25	61N	3.582	25C25	61CG	3.623
25C25	650	4.058	25C25	60C	4.074	25C25	61CA	4.084
25C25	590	4.120	25C25	60CA	4.277	25C25	610D1	4.827
25C25	65C	4.828	25C25	600	4.832	25C25	66CA	4.907
25C26	61CB	3.331	25C26	650	3.342	25C26	610D2	3.552
25C26	61CG	3.818	25C26	61N	3.860	25C26	65C	3.890
25C26	61CA	4.226	25C26	640	4.287	25C26	66N	4.362
25C26	66CA	4.392	25C26	60C	4.694	25C26	65CA	4.697
25C26	64C	4.855	25C26	610D1	4.925	25C26	60CA	4.944
25C27	650	3.537	25C27	65C	3.661	25C27	66N	3.764
25C27	66CA	3.789	25C27	67CE2	4.460	25C27	65CA	4.463
25C27	640	4.501	25C27	61CB	4.521	25C27	66C	4.533
25C27	61N	4.713	25C27	67CD2	4.833	25C27	67CZ	4.873
25C27	610D2	4.935	25C27	660	4.986	25C28	67CE2	3.131
25C28	67CD2	3.493	25C28	67CZ	3.739	25C28	66CA	3.757
25C28	670H	4.121	25C28	66C	4.151	25C28	66N	4.180
25C28	67CG	4.339	25C28	650	4.368	25C28	67N	4.383
25C28	65C	4.453	25C28	67CE1	4.537	25C28	60ND2	4.576
25C28	660	4.719	25C28	67CD1	4.789	25C29	67CE2	3.099
25C29	67CD2	3.367	25C29	60ND2	4.065	25C29	590	4.096
25C29	67CZ	4.107	25C29	66CA	4.348	25C29	67CG	4.510
25C29	670H	4.532	25C29	67N	4.689	25C29	60CA	4.697
25C29	66C	4.785	25C29	700D1	4.794	25C29	650	4.942
25C30	65C	3.364	25C30	66N	3.477	25C30	650	3.676
25C30	65CA	3.705	25C30	640	3.715	25C30	66CA	3.996
25C30	64C	4.521	25C30	65N	4.571	25C30	66C	4.606
25C30	660	4.690	25031	66N	4.304	25031	65C	4.471
25031	670H	4.497	25031	67CZ	4.562	25031	65CA	4.615
25031	640	4.643	25031	66CA	4.801	25031	660	4.804
25031	67CE1	4.862	25031	67CE2	4.899	25031	650	4.995
25031	66C	4.998	25C32	660	4.210	25C32	66N	4.247
25C32	65CA	4.548	25C32	55C	4.675	25C32	67CE1	
25C32	66C	4.745	25C32	67CZ	4.797	25C32	66CA	4.862
25C32		4.906	25033	67CE1		25033	67CZ	4.445
25033	670H	4.485	25033	660	4.628			4.718
25033	2530H2	4.812	25033	1600	4.869	25C34	660	3.728

## TABLE XXV

25C34	66N	4.086	25C34	65CA	4.283	25C34	65C	4.685
25C34	25 <i>S</i> G	4.701	25C34	66C	4.710	25C34	1610	4.715
25C34	161C	4.780	25C35	660	3.094	25C35	66C	4.237
25C35	68CE	4.248	25C35	6 <b>6N</b>	4.449	25C35	163CB	4.779
25C35	163N	4.929	25C35	66CA	4.993	25C36	660	4.248
25C36	134CB	4.455	25C36	209CD2	4.608	25C36	68CE	4.627
25C36	1600	4.689	25C36	162N	4.700	25C36	160C	4.804
25C36	161C	4.832	25C36	161CA	4.852	25C36	67CD1	4.855
25C36	67CE1	4.866	25C36	161N	4.905	25C36	160CB	4.964
25C37	162N	3.453	25C37	134CB	3.566	25C37	162C	3.722
25C37	163N	3.858	25C37	1620	3.881	25C37	161C	3.888
25C37	161N	4.019	25C37	162CA	4.040	25C37	161CA	4.109
25C37	160C	4.197	25C37	163CA	4.335	25C37	163CB	4.414
25C37	1600	4.461	25C37	160CB	4.479	25C37	134CA	4.485
25C37	1610	4.643	25C37	160CA	4.832	25C37	68CE	4.842
25C37	209CD2	4.996	25C38	209CD2	3.377	25C38	67CD1	3.746
25C38	67CE1	3.972	25C38	68CE	4.019	25C38	134CB	4.063
25C38	660	4.117	25C38	209CG	4.387	25C38	67CG	4.813
25C38	67CA	4.927	25C39	65CA	3.495	25C39	25 <b>S</b> G	3.606
25C39	66N	3.612	25C39	660	3.741	25C39	26CD1	4.029
25C39	65C	4.099	25C39	26CB	4.554	25C39	26CG	4.652
25C39	66C	4.666	25C39	65N	4.740	25C39	26N	4.747
25C39	66CA	4.755	25C39	230	4.898	25C39	1610	4.941
25040	66N	2.789	25040	660	2.930	25040	26CD1	2.937
25040	65CA	3.211	25040	26CB	3.444	25040	26CG	3.455
25040	65C	3.479	25040	66C	3.767	25040	66CA	3.842
25040	25 <b>S</b> G	3.892	25040	26N	3.999	25040	26NE1	4.011
25040	26CA	4.273	25040	65N	4.534	25040	230	4.581
25040	26CD2	4.697	25040	650	4.703	25040	68CE	4.926
25040	26CE2	4.943	25040	25N	4.988	25N41	25SG	2.879
25N41	65CA	3.789	25N41	1610	4.253	25N41	230	4.382
25N41	26CD1	4.417	25N41	66N	4.577	25N41	25CB	4.611
25N41	23C	4.734	25N41	65N	4.734	25N41	65C	4.774
25N41	25N	4.800	25N41		4.855	25N41	26 <b>N</b>	4.870
		4.874		163N				
25C42	230	3.257		23C	3.448			3.529
25C42		3.714		65CA	3.771		25CB	3.843
25C42		3.956	25C42		3.966		26N	4.147
25C42		4.156	25C42		4.282	25C42		4.390
25C42	65N	4.469	25C42	26NE1	4.524			4.662
25C42	26CG	4.728	25C42	66N	4.771	25C42	65C	4.825
25C42		4.981		55N	3.501	25N43	65CA	3.665
25N43	660	3.722	25N43	65C	3.917	25N43	66C	4.394

TABLE XXV

25N43 66CA 4.413 25N43 640 4.690 25N43 650 4.923

25N43 65N 4.930

#### TABLE XXVI

Table of distances in Ångstroms between atoms of the inhibitor and protein for all protein atoms within 5 Ångstroms of the inhibitor 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	2420H2	3.241	25C1	180D1	3.446	25C1	184CD1	3.907
25C1	184CB	3.929	25C1	184CG	3.941	25C1	184CA	4.202
25C1	18CG	4.369	25C1	18ND2	4.563	25C1	184NE1	4.640
25C1	1840	4.676	25C1	184CD2	4.729	25C1	184C	4.752
25C1	21NE2	4.852	25C1	20N	4.956	25C2	184CD1	3.751
25C2	184CG	3.778	25C2	184NE1	4.100	25C2	184CD2	4.156
25C2	184CB	4.179	25C2	184CE2	4.339	25C2	2420H2	4.411
25C2	180D1	4.705	25C2	184CA	4.841	25C2	184CE3	4.910
25C3	184NE1	3.789	25C3	184CD1	3.817	25C3	184CE2	4.203
25C3	184CG	4.246	25C3	184CD2	4.475	25C3	200	4.505
25C3	184CZ2	4.908	25C3	21NE2	4.969	25C4	200	3.140
25C4	19CG	4.004	25C4	20C	4.011	25C4	184CD1	4.042
25C4	184NE1	4.079	25C4	21NE2	4.367	25C4	20N	4.403
25C4	19CD	4.509	25C4	20CA	4.543	25C4	210E1	4.778
25C4	184CG	4.809	25C4	180D1	4.811	25C4	184CE2	4.867
25C4	21CD	4.895	25C4	19NE2	4.938	25C4	21N	4.965
25C5	200	2.695	25C5	20N	3.202	25C5	20C	3.263
25C5	20CA	3.407	25C5	180D1	3.589	25C5	19CG	3.659
25C5	21NE2	3.950	25C5	19C	4.127	25C5	184CD1	4.183
25C5	21N	4.369	25C5	2420H2	4.392	25C5	18CG	4.548
25C5	19CD	4.599	25C5	184NE1	4.619	25C5	19N	4.620
25C5	19CA	4.668	25C5	19CB	4.730	25C5	21CD	4.753
25C5	18ND2	4.801	25C5	190	4.927	25 <b>C</b> 5	184CG	4.931
25C5	1830	4.975	25C5	210E1	4.987	25C6	180D1	2.695
25C6	2420H2	3.232	25C6	20N	3.572	25C6	18CG	3.627
25C6	18ND2	3.837	25C6	20CA	3.842	25C6	200	3.869
25C6	184CD1	4.122	25C6	20C	4.133	25C6	21NE2	4.219
25 <b>C</b> 6	19CG	4.448	25C6	184CA	4.454	25C6	184CG	4.538
25C6	184CB	4.595	25C6	19N	4.613	25C6	19C	4.668

## TABLE XXVI

25C6	1830	4.825	25C6	2440H2	4.841	25C6	184NE1	4.878
25C7	200	2.832	25C7	19CG	3.901	25C7	20C	3.960
25C7	19CD	4.075	25C7	19NE2	4.108	25C7	220	4.387
25C7	184NE1	4.479	25C7	22N	4.658	25C7	21CA	4.703
25C7	190E1	4.733	25C7	210E1	4.734	25C7	21N	4.779
25C7	184CD1	4.791	25C7	20N	4.797	25C7	21NE2	4.850
25C7	20CA	4.870	2508	200	4.185	2508	184NE1	4.318
2508	19NE2	4.505	2508	19CD	4.575	2508	19CG	4.801
2508	184CE2	4.904	2508	184CD1	4.980	2508	190E1	4.989
25C9	184NE1	3.701	25C9	19NE2	4.091	25C9	19CD	4.179
25C9	184CE2	4.209	25C9	184CZ2	4.214	25C9	190E1	4.322
25C9	162CE1	4.669	25C9	184CD1	4.680	25C9	19CG	4.819
25010	184NE1	2.553	25010	184CE2	3.148	25010	184CZ2	3.297
25010	190E1	3.438	25010	19CD	3.591	25010	184CD1	3.650
25010	162CE1	3.653	25010	19NE2	3.822		162ND1	4.295
25010	19CG	4.332	25010	184CD2	4.379	25010	162NE2	4.393
25010	184CH2	4.579	25010	184CG	4.611		162ND1	3.950
25C11	162CE1	4.196	25C11	184CZ2	4.648	25C11	19NE2	4.924
25C11	162CG	4.968	25C12	162ND1	4.717	25C12	1610	4.982
25C13	1610D1	4.017	25C13	162ND1	4.533	25C13	161CG	4.712
25C13	1610	4.749	25C13	184CZ2	4.831	25C13	162CB	4.860
25C13	162CG	4.864	25C13	1370	4.980		1610D1	3.145
25C14	162ND1	3.354	25C14	162CB	3.366	25C14	162CG	3.508
25C14	1610	3.685	25C14	161C	3.990		162CA	4.004
25C14	161CG	4.037	25C14	162N	4.168	25C14	162CE1	4.261
25C14	137CB	4.367	25C14	161CB	4.378		162CD2	4.476
25C14	184CZ2	4.641	25C14	137C	4.815	25C14	162NE2	4.847
25C14	161CA	4.867	25C14	1370	4.907		137CA	5.000
25C15	1370	4.063	25C15	184CZ2	4.064		184CH2	4.220
25C15	137C	4.530		161001			138CA	4.806
25C15	138N	4.887	25C15	137CB	4.908			
25C16	25SG	4.159	25C16	19NE2			162CE1	
25C16	23CA	4.579		1610			23CA	3.462
25017	19NE2	3.921		23C				4.429
25017	220	4.568	25017	25SG	4.579		22C	4.858
25017	1.9CD	4.877	25017	230	4.921			4.983
25N18	25SG	3.257	25N18	162ND1	3.402	25N18	1610	3.468

## TABLE XXVI

25N18	162CE1	4.081	25N18 25CB	4.457	25N18 162CG	4.523
25N18	161C	4.569	25N18 162CA	4.582	25N18 162CB	4.845
25N18	19NE2	4.859	25C19 25SG	2.794	25C19 1610	3.684
25C19	25CB	4.234	25C19 162ND1	4.247	25C19 23CA	4.366
25C19	230	4.457	25C19 23C	4.497	25C19 25N	4.673
25C19	161C	4.876	25C19 162CE1	4.889	25C19 19NE2	4.898
25N20	19NE2	4.613	25N20 184NE1	4.773	25N20 184CZ2	4.805
25N20	162CE1	4.936	25N20 19CD	4.991	25C21 1610	2.879
25C21	25SG	3.218	25C21 161C	4.048	25C21 162CA	4.631
25C21	162N	4.811	25C21 65CA	4.831	25C21 162ND1	4.843
25C21	25CB	4.939	25C21 163N	4.976	25C22 25SG	1.746
25C22	25CB	2.980	25C22 25N	3.156	25C22 25CA	3.592
25C22	230	3.702	25C22 23C	3.712	25C22 23CA	4.038
25C22	19NE2	4.077	25C22 24N	4.130	25C22 26N	4.232
25C22	162ND1	4.246	25C22 24C	4.298	25C22 25C	4.374
25C22	162CE1	4.608	25C22 24CA	4.620	25C22 1610	4.649
25C22	26CD1	4.828	25C22 19OE1	4.874	25C22 19CD	4.942
25023	25SG	2.426	25023 25N	2.644	25023 25CB	2.944
25023	19NE2	2.944	25023 23C	2.957	25023 24N	3.189
25023	23CA	3.211	25023 230	3.326	25023 25CA	3.349
25023	24C	3.666	25023 24CA	3.882	25023 19CD	3.941
25023	190E1	4.099	25023 220	4.365	25023 25C	4.451
25023	26N	4.481	25023 23N	4.525	25023 162ND	
25023	162CE1	4.629	25023 240	4.773	25023 22C	4.910
25C24	65CA	4.095	25C24 66N	4.264	25C24 65C	4.356
25C24	640	4.547	25C24 660	4.628	25C25 66O	3.688
25C25	1610	3.862	25C25 66N	4.313	25C25 161C	4.458
25C25	66C	4.741	25C25 65CA	4.778	25C25 161CA	
25C25	25 <i>S</i> G	4.963	<b>25C25</b> 65C	4.974	25C26 66O	3.481
25C26	1610	4.269	25C26 161C	4.491	25C26 163N	4.695
25C26	66C	4.696	25C26 160C			
25C26	162N	4.915	25C26 161CA			
25C27	160C	4.089	25C27 161CA			
	161N		25C27 1610			
25C27	160CB	4.699	25C27 134CB	4.734	25C27 162N	4.770
25C27	209CD2	4.918	25C28 160C	3.351	25C28 160C	
25C28	67CE1	4.588	25C28 161CA	4.739	25C28 66O	4.818

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25C28	161N	4.934	25C28	67CD1	4.955	25C29	209CD2	3.439
25C29	134CB	3.686	25C29	1600	4.049	25C29	160CB	4.234
25C29	160C	4.490	25C29	67CD1	4.759	25C29	660	4.795
25C29	209CG	4.839	25C29	67CE1	4.896	25C30	660	3.533
25C30	66N	3.616	25C30	25SG	3.883	25C30	65CA	3.972
25C30	26CD1	4.246	25C30	26CB	4.295	25C30	65C	4.296
25C30	1610	4.302	25C30	66C	4.432	25C30	66CA	4.591
25C30	26CG	4.623	25C30	26N	4.737	25C30	163CB	4.852
25C30	163N	4.940	25031	660	2.635	25031	66N	2.846
25031	26CB	3.199	25031	26CD1	3.340	25031	66C	3.423
25031	26CG	3.516	25031	66CA	3.653	25031	65C	3.792
25031	65CA	3.821	25031	26N	4.122	25031	26CA	4.160
25031	25 <i>S</i> G	4.278	25031	26NE1	4.531	25031	67N	4.618
25031	163CB	4.642	25031	26CD2	4.797	25031	650	4.959
25N32	25SG	2.963	25N32	1610	3.860	25N32	65CA	4.049
25N32	66N	4.382	25N32	230	4.384	25N32	26CD1	4.398
25N32	26N	4.601	25 <b>N3</b> 2	25CB	4.628	25N32	25N	4.764
25N32	65C	4.790	25N32	660	4.806	25N32	163N	4.815
25N32	26CB	4.836	25N32	161C	4.880	25N32	23C	4.985
25C33	25 <b>S</b> G	2.418	25C33	230	3.256	25C33	25N	3.403
25C33	26CD1	3.587	25C33	26N	3.672	25C33	25CB	3.751
25C33	23C	3.758	25C33	25CA	3.959	25C33	65CA	3.999
25C33	25C	4.264	25C33	24N	4.334	25C33	24C	4.346
25C33	26CG	4.388	25C33	26CB	4.392	25C33	23CA	4.425
25C33	66N	4.510	25C33	24CA	4.531	25C33	26CA	4.616
25C33	26NE1	4.621	25C33	65C	4.831	25C33	65N	4.901
25C33	1610	4.934	25N34	660	3.486	25N34	6 <b>6N</b>	3.771
25N34	65CA	4.261	25N34	65C	4.264	25N34	6 <b>6</b> C	4.330
25N34	66CA	4.464	25N34	1610	4.991			

#### **TABLE XXVII**

Table of distances in Angstroms between atoms of the inhibitor and protein for all protein atoms within 5 Angstroms of the inhibitor 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N[N-(methyl)-L-leucyl)]-3-pyrrolidinone.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	2420H2	3.241	25C1	180D1	3.446	25C1	184CD1	3.907
25C1	184CB	3.929	25C1	184CG	3.941	25C1	184CA	4.202
25C1	18CG	4.369	25C1	18ND2	4.563	25C1	184NE1	4.640
25C1	1840	4.676	25C1	184CD2	4.729	25C1	184C	4.752
25C1	21NE2	4.852	25C1	20N	4.956	25C2	184CD1	3.751
25C2	184CG	3.778	25C2	184NE1	4.100	25C2	184CD2	4.156
25C2	184CB	4.179	25C2	184CE2	4.339	25C2	2420H2	4.411
25C2	180D1	4.705	25C2	184CA	4.841	25C2	184CE3	4.910
25C3	184NE1	3.789	25C3	184CD1	3.817	25C3	184CE2	4.203
25C3	184CG	4.246	25C3	184CD2	4.475	25C3	200	4.505
25C3	184CZ2	4.908	25C3	21NE2	4.969	25C4	200	3.140
25C4	19CG	4.004	25C4	20C	4.011	25C4	184CD1	4.042
25C4	184NE1	4.079	25C4	21NE2	4.367	25C4	20N	4.403
25C4	19CD	4.509	25C4	20CA	4.543	25C4	210E1	4.778
25C4	184CG	4.809	25C4	180D1	4.811	25C4	184CE2	4.867
25C4	21CD	4.895	25C4	19NE2	4.938	25C4	21N	4.965
25C5	200	2.695	25C5	20 <b>N</b>	3.202	25C5	20C	3.263
25C5	20CA	3.407	25C5	180D1	3.589	25C5	19CG	3.659
25C5	21NE2	3.950	2505	19C	4.127	25C5	184CD1	4.183
25C5	21N	4.369	25C5	2420H2	4.392	25C5	18CG	4.548
25C5	19CD	4.599	25C5	184NE1	4.619	25C5	19N	4.620
25C5	19CA	4.668	25C5	19CB	4.730	25C5	21CD	4.753
25C5	19ND2	4.801	25C5	190	4.927	25C5	184CG	4.931
25C5	1830	4.975	25C5	210E1	4.987	25C6	180D1	2.695
25C6	2420H2	3.232	25C6	20N	3.572	25 <b>C</b> 6	18CG	3.627
25C6	18ND2	3.837	25€€	20CA	3.842	25C6	200	3.869
25C6	184CD1	4.122	25C6	20C	4.133	25C6	21NE2	4.219
25C6	19CG	4.448	25C6	184CA	4.454	25C6	184CG	4.538
25C6	184CB	4.595	25C6	19N	4.613	25C6	19C	4.668

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25C6	1830	4.825	25C6	2440H2	4.841	25C6	184NE1	4.878
25C7	200	2.832	25C7	19CG	3.901	25C7	20C	3.960
25C7	19CD	4.075	25C7	19NE2	4.108	25C7	220	4.387
25C7	184NE1	4.479	25C7	22N	4.658	25C7	21CA	4.703
25C7	190E1	4.733	25C7	210E1	4.734	25C7	21N	4.779
25C7	184CD1	4.791	25C7	20N	4.797	25C7	21NE2	4.850
25C7	20CA	4.870	2508	200	4.185	2508	184NE1	4.318
2508	19NE2	4.505	2508	19CD	4.575	2508	19CG	4.801
2508	184CE2	4.904	2508	184CD1	4.980	2508	190E1	4.989
25C9	184NE1	3.701	25 <b>C</b> 9	19NE2	4.091	25C9	19CD	4.179
25C9	184CE2	4.209	25C9	184CZ2	4.214	25C9	190E1	4.322
25C9	162CE1	4.669	25C9	184CD1	4.680	25C9	19CG	4.819
25010	184NE1	2.553	25010	184CE2	3.148	25010	184CZ2	3.297
25010	190E1	3.438	25010	19CD	3.591	25010	184CD1	3.650
25010	162CE1	3.653	25010	19NE2	3.822	25010	162ND1	4.295
25010	19CG	4.332	25010	184CD2	4.379	25010	162NE2	4.393
25010	184CH2	4.579	25010	184CG	4.611	25C11	162ND1	3.950
25C11	162CE1	4.196	25C11	184CZ2	4.648	25C11	19NE2	4.924
25C11	162CG	4.968	25C12	162ND1	4.717	25C12	1610	4.982
25C13	1610D1	4.017	25C13	162ND1	4.533	25C13	161CG	4.712
25C13	1610	4.749	25C13	184CZ2	4.831	25C13	162CB	4.860
25C13	162CG	4.864	25C13	1370	4.980	25C14	1610D1	3.145
25C14	162ND1	3.354	25C14	162CB	3.366	25C14	162CG	3.508
25C14	1610	3.685	25C14	161C	3.990	25C14	162CA	4.004
25C14	161CG	4.037	25C14	162N	4.168	25C14	162CE1	4.261
25C14	137CB	4.367	25C14	161CB	4.378	25C14	162CD2	4.476
25C14	184CZ2	4.641	25C14	137C	4.815	25C14	162NE2	4.847
25C14	161CA	4.867	25C14	1370	4.907	25C14	137CA	5.000
25C15		4.063		184CZ2	4.064		184CH2	4.220
25C15	137C	4.530		1610D1				
25C15	138N	4.887		137CE			162ND1	
25C16				19NE2				4.375
25C16	23CA	4.579	25C16	1610				
25017	19NE2	3.921	25017	23C	4.371	25017	23N	4.429
25017	220	4.568	25017	25SG				4.858
25017	19CD	4.877	25017	230	4.921	25017	24N	4.983
25N18	25 <b>S</b> G	3.257	25N18	162ND1	3.402	25N18	1610	3.468

25N18	162CE1	4.081	25N18	25CB	4.457	25N18	162CG	4.523
25N18	161C	4.569	25N18	162CA	4.582	25N18	162CB	4.845
25N18	19NE2	4.859	25C19	25SG	2.794	25C19	1610	3.684
25C19	25CB	4.234	25C19	162ND1	4.247	25C19	23CA	4.366
25C19	230	4.457	25C19	23C	4.497	25C19	25N	4.673
25C19	161C	4.876	25C19	162CE1	4.889	25C19	19NE2	4.898
25N20	19NE2	4.613	25N20	184NE1	4.773	25N20	184CZ2	4.805
25N20	162CE1	4.936	25N20	19CD	4.991	25C21	1610	2.879
25C21	25SG	3.218	25C21	161C	4.048	25C21	162CA	4.631
25C21	162N	4.811	25C21	65CA	4.831	25C21	162ND1	4.843
25C21	25CB	4.939	25C21	163N	4.976	25C22	25 <i>S</i> G	1.746
25C22	25CB	2.980	25C22	25N	3.156	25C22	25CA	3.592
25C22	230	3.702	25C22	23C	3.712	25C22	23CA	4.038
25C22	19NE2	4.077	25C22	24N	4.130	25C22	26N	4.232
25C22	162ND1	4.246	25C22	24C	4.298	25C22	25C	4.374
25C22	162CE1	4.608	25C22	24CA	4.620	25C22	1610	4.649
25C22	26CD1	4.828	25C22	190E1	4.874	25C22	19CD	4.942
25023	25 <b>S</b> G	2.426	25023	25N	2.644	25023	25CB	2.944
25023	19NE2	2.944	25023	23C	2.957	25023	24N	3.189
25023	23CA	3.211	25023	230	3.326	25023	25CA	3.349
25023	24C	3.666	25023	24CA	3.882	25023	19CD	3.941
25023	190E1	4.099	25023	220	4.365	25023	25C	4.451
25023	26N	4.481	25023	23N	4.525	25023	162ND1	4.590
25023	162CE1	4.629	25023	240	4.773	25023	22C	4.910
25C24	65CA	4.095	25C24	6 6 N	4.264	25C24	65C	4.356
25C24	640	4.547	25C24	660	4.628	25C25	660	3.688
25C25	1610	3.862	25C25	66N	4.313	25C25	161C	4.458
25C25	66C	4.741	25C25	65CA	4.778	25C25	161CA	4.939
25C25	25SG	4.963	25C25	65C	4.974	25C26	660	3.481
25C26	1610		25C26		4.491	25C26		4.695
25C26	66C	4.696	25C26	1600	4.785			
25C26	162N	4.915	25C26	161CA	4.923	25C27	1600	3.472
25C27	160C	4.089	25C27	161CA	4.307	25C27	161C	4.320
25C27	161N	4.440	25C27	1610	4.452	25C27	660	4.548
25C27	160CB	4.699	25C27	134CB	4.734		162N	4.770
25C27	209CD2	4.918	25028	1600	3.351		160C	4.338
25C28	67CE1	4.588	25C28	151CA	4.739	25C28	660	4.818

25C28	161N	4.934	25C28	67CD1	4.955	25C29	209CD2	3.439
25C29	134CB	3.686	25C29	1600	4.049	25C29	160CB	4.234
25C29	160C	4.490	25C29	67CD1	4.759	25C29	660	4.795
25C29	209CG	4.839	25C29	67CE1	4.896	25C30	660	3.533
25C30	66N	3.616	25C30	25SG	3.883	25C30	65CA	3.972
25C30	26CD1	4.246	25C30	26CB	4.295	25C30	65C	4.296
25C30	1610	4.302	25C30	66C	4.432	25C30	66CA	4.591
25C30	26CG	4.623	25C30	26N	4.737	25C30	163CB	4.852
25C30	163N	4.940	25031	660	2.635	25031	66N	2.846
25031	26CB	3.199	25031	26CD1	3.340	25031	66C	3.423
25031	26CG	3.516	25031	66CA	3.653	25031	65C	3.792
25031	65CA	3.821	25031	26N	4.122	25031	26CA	4.160
25031	25SG	4.278	25031	26NE1	4.531	25031	67N	4.618
25031	163CB	4.642	25031	26CD2	4.797	25031	650	4.959
25N32	25 <b>S</b> G	2.963	25N32	1610	3.860	25N32	65CA	4.049
25N32	66N	4.382	25N32	230	4.384	25N32	26CD1	4.398
25N32	26N	4.601	25N32	25CB	4.628	25N32	25N	4.764
25N32	65C	4.790	25N32	660	4.806	25N32	163N	4.815
25N32	26CB	4.836	25N32	161C	4.880	25N32	23C	4.985
25C33	25SG	2.418	25C33	230	3.256	25C33	25N	3.403
25C33	26CD1	3.587	25C33	26N	3.672	25C33	25CB	3.751
25C33	23C	3.758	25C33	25CA	3.959	25C33	65CA	3.999
25C33	25C	4.264	25C33	24N	4.334	25C33	24C	4.346
25C33	26CG	4.388	25C33	26CB	4.392	25C33	23CA	4.425
25C33	66N	4.510	25C33	24CA	4.531	25C33	26CA	4.616
25C33	26NE1	4.621	25C33	65C	4.831	25C33	65N	4.901
25C33	1610	4.934	25N34	660	3.486	25N34	66N	3.771
25N34	65CA	4.261	25N34	65C	4.264	25N34	6 <b>6</b> C	4.330
25N34	66CA	4.464	25N34	1610	4.991			

#### **TABLE XXVIII**

Table of distances in Angstroms between atoms of the inhibitor and protein for all protein atoms within 5 Angstroms of the inhibitor 1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one.

Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.	Atom	1 Atom 2	Dist.
25C1	200	2.824	25C1	20C	3.956	25C1	184NE1	4.078
25C1	184CD1	4.178	25C1	21CA	4.600	25C1	20N	4.704
25C1	21N	4.706	25C1	21NE2	4.738	25C1	19CB	4.859
25C1	19CD	4.929	25C1	20CA	4.947	25C1	184CE2	4.949
25C1	19CG	4.998	25C2	200	3.367	25C2	21NE2	4.119
25C2	2410H2	4.309	25C2	20C	4.592	25C2	184NE1	4.819
25C2	184CD1	4.860	25C3	2410H2	3.093	25C3	200	3.775
25C3	21NE2	4.450	25C3	184CD1	4.878	25C3	20C	4.988
25C4	2410H2	3.480	25C4	200	3.717	25C4	180D1	3.947
25C4	184CD1	4.219	25C4	184CG	4.493	25C4	1840	4.549
25C4	184CB	4.602	25C4	184CA	4.747	25C4	18CG	4.754
25C4	18ND2	4.754	25C4	20C	4.826	25C4	184NE1	4.840
25C4	20N	4.964	25C4	184C	4.980	25C5	200	3.239
25C5	180D1	3.271	25C5	184CD1	3.400	25C5	20N	3.933
25C5	184CG	4.002	25C5	184NE1	4.111	25C5	184CA	4.118
25C5	20C	4.232	25C5	18CG	4.245	25C5	184CB	4.256
25C5	19CB	4.402	25C5	1830	4.499	25C5	1840	4.503
25C5	20CA	4.510	25C5	19N	4.552	25C5	184C	4.587
25C5	18ND2	4.606	25C5	19C	4.765	25C5	19CA	4.817
25C5	2410H2	4.867	25C5	184CD2	4.968	25C6	200	2.740
25C6	184CD1	3.361	25C6	184NE1	3.660	25C6	20C	3.746
25C6	20N	3.761	25C6	19CB	3.788	25C6	180D1	4.136
25C6	19CG	4.290	25C6	184CG	4.321	25C6	20CA	4.335
25C6	19C	4.361	25C6	19CD	4.428	25C6	19CA	4.488
25C6	190E1	4.526	25C6	19N	4.653	25C6	184CE2	4.729
25C6	1830	4.737	25C6	21N	4.769	25C6	184CA	4.918
25C6	184CB	4.981	25C6	19NE2	4.990	2507	200	3.287
2507	21CA	4.138	2507	20C	4.173	2507	19NE2	4.269
2507	21C	4.370	2507	1.9CD	4.416	2507	184NE1	4.459

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2507	210	4.466	2507	220	4.540	2507	21N	4.558
2507	19CG	4.663	2507	190E1	4.850	2507	184CD1	4.897
2507	19CB	4.977	25C8	19NE2	4.003	25C8	184NE1	4.235
25C8	19CD	4.404	25C8	200	4.665	25C8	190E1	4.715
25C8	220	4.861	25C8	210	4.937	25C8	184CE2	4.948
25C9	184NE1	3.199	25C9	19NE2	3.212	25C9	19CD	3.472
25C9	190E1	3.533	25C9	184CE2	4.000	25C9	184CD1	4.152
25C9	184CZ2	4.206	25C9	19CG	4.414	25C9	162NE2	4.653
25C9	220	4.860	25C9	162CD2	4.867	25C9	19CB	4.910
25C10	184NE1	3.388	25C10	184CZ2	3.605	25C10	19 <b>NE</b> 2	3.666
25C10	184CE2	3.828	25C10	162CD2	3.878	25C10	190E1	3.887
25C10	162NE2	3.948	25C10	19CD	4.067	25C10	184CD1	4.615
25C10	162CG	4.799	25C10	184CH2	4.859	25C10	162CE1	4.873
25C11	184CZ2	4.084	25C11	184NE1	4.519	25C11	162CD2	4.567
25C11	184CE2	4.664	25C11	19NE2	4.718	25C11	162NE2	4.920
25C13	19NE2	4.978	25\$14	162CD2	3.947	25S14	184CZ2	4.113
25 <b>S</b> 14	162CG	4.222	25\$14	162CB	4.332	25S14	1610	4.574
25S14	162NE2	4.657	25S14	1610D1	4.760	25 <b>S</b> 14	184CH2	4.865
25015	184CZ2	3.079	25015	184CH2	3.580	25015	162CG	3.856
25015	162CD2	3.881	25015	1370	3.985	25015	162CB	4.051
25015	184CE2	4.275	25015	137CB	4.300	25015	162ND1	4.343
25015	162NE2	4.364	25015	1610D1	4.610	25015	162CE1	4.620
25015	137C	4.663	25015	184NE1	4.807	25015	184CZ3	4.973
25016	1610D1	4.484	25016	1610	4.575	25016	161CG	4.963
25N17	162CD2	2.828	25 <b>N1</b> 7	162CG	3.217	25N17	1610	3.256
25N17	162CB	3.276	25 <b>N</b> 17	162CA	3.672	25N17	162NE2	3.893
25N17	161C	4.006	25N17	25 <i>S</i> G	4.125	25N17	162N	4.251
25N17	162ND1	4.353	25N17	1610D1	4.593	25N17		4.665
25N17	184CZ2	4.778	25N17	25CB	4.819	25C18	1610	3.031
25C18		3.290		162CD2	3.499		162CA	4.069
25C18	161C	4.107	25C18	162CG	4.151	25C18	162CB	4.188
25C18	25CE	4.359	25018	19NE2	4.573	25C18	162NE2	4.594
25C18	162N	4.597	25C19	25SG	1.931	25C19	25CB	3.158
25C19	162CD2	3.616	25C19	1610	3.716	25C19	19NE2	3.740
25C19	25N	4.227	25C19	25CA	4.306	25C19	23CA	4.380
25C19	162CA	4.405	25C1.9	230	4.494		23C	
25C19	162CG	4.541	25C19	1.62NE2	4.577	25C19	19CD	4.643

25C19	190E1	4.705	25C19	161C	4.813	25C19	162CB	4.819
25020	19NE2	2.778	25020	25SG	2.890	25020	23CA	3.251
25020	25CB	3.629	25020	23C	3.723	25020	19CD	3.918
25020	230	4.030	25020	25N	4.152	25020	190E1	4.290
25020	24N	4.342	25020	162CD2	4.356	25020	23N	4.524
25020	25CA	4.566	25020	220	4.681	25020	1610	4.912
25C21	25SG	2.532	25C21	1610	3.420	25C21	230	4.107
25C21	25CB	4.220	25C21	23C	4.540	25C21	65CA	4.542
25C21	161C	4.578	25C21	23CA	4.599	25C21	162CA	4.793
25C21	25N	4.822	25C21	162CD2	4.980	25N22	1610	2.689
25N22	25SG	2.783	25N22	161C	3.649	25N22	162CA	4.003
25N22	162N	4.221	25N22	163N	4.453	25N22	25CB	4.578
25N22	162C	4.691	25 <b>N2</b> 2	161CA	4.693	25N22	65CA	4.839
25C23	1610	3.461	25C23	25 <b>S</b> G	3.771	25C23	161C	4.205
25C23	660	4.217	25C23	65CA	4.220	25C23	66N	4.268
25C23	162N	4.828	25C23	162CA	4.844	25C23	65C	4.845
25C23	26CD1	4.890	25C23	161CA	4.937	25C23	26CB	4.974
25024	65CA	3.024	25024	66N	3.208	25024	65C	3.657
25024	660	3.830	25024	65N	4.287	25024	26CD1	4.326
25024	25 <b>S</b> G	4.394	25024	66CA	4.437	25024	1610	4.526
25024	66C	4.595	25024	640	4.818	25024	230	4.843
25024	26CG	4.879	25024	26CB	4.888	25024	650	4.923
25024	64C	4.972	25C25	1610	3.652	25C25	660	3.880
25C25	161C	3.949	25C25	162N	4.375	25C25	161CA	4.493
25C25	163N	4.584	25C25	25SG	4.592	25C25	162CA	4.608
25C25	162C	4.759	25C25	66N	4.805	25C25	163CB	4.922
25C26	660	3.186	25C26	163CB	3.703	25C26	26CB	3.850
25C26	163N	4.052	25C26	25SG	4.357	25C26	163CA	4.390
25C26	66C	4.421	25C26	26CA	4.561	25C26	162C	4.598
25C26	68SD	4.628	25C26	1610	4.666	25C26	6 <b>6N</b>	4.721
25C26	26N	4.729	25C26	26CG	4.770	25C26	162CA	4.869
25C26	161C	4.907	25C26	162N	4.970	25C26	26CD1	4.999
25C27	163CB	3.337	25C27	68SD	3.583	25C27	660	3.590
25C27	163CA	4.029	25C27	163N	4.096	25C27	68CE	4.156
25C27	134CB	4.190	25C27	209CD2	4.248	25C27	26CB	4.361
25C27	162C	4.597	25C27	66C	4.797			4.893
25C27	26CA	4.943	25C27	67CA	4.946	25C28	134CB	3.049

25C28	163CB	3.699	25C28	163N	3.714	25C28	162C	3.789
25C28	1620	3.807	25C28	163CA	3.820	25C28	209CD2	3.901
25C28	134CA	3.967	25C28	162N	4.239	25C28	162CA	4.478
25C28	68CE	4.498	25C28	68SD	4.535	25C28	161C	4.582
25C28	660	4.956	25C28	134C	4.964	25C28	1610	4.966
25C28	161N	4.992	25C29	660	2.949	25C29	209CD2	3.567
25C29	68SD	3.617	25C29	67CA	3.759	25C29	67CD1	3.963
25C29	66C	3.998	25C29	68CE	4.231	25C29	67N	4.370
25C29	68N	4.410	25C29	67C	4.555	25C29	67CB	4.556
25C29	67CG	4.589	25C29	163CB	4.636	25C29	134CB	4.685
25C29	2340H2	4.722	25C29	26CB	4.726	25C29	67CE1	4.755
25N30	660	3.644	25N30	66N	4.623	25N30	1610	4.640
25 <b>N</b> 30	66C	4.640	25N30	161C	4.793	25N30	161CA	4.888
25N30	1600	4.908	25C31	1600	3.792	25C31	161CA	4.412
25C31	160C	4.476	25C31	161C	4.719	25C31	660	4.743
25C31	161N	4.777	25C31	1610	4.849	25C31	67CE1	4.934
25032	1600	2.720	25032	160C	3.270	25032	161CA	3.397
25032	161N	3.577	25032	161C	3.869	25032	160CB	4.202
25032	1610	4.268	25032	160CA	4.363	25032	162N	4.395
25032	161CB	4.698	25C33	67CE1	3.801	25C33	1600	4.307
25C33	67CZ	4.397	25C33	67CD1	4.397	25C33	670H	4.480
25C34	1600	3.849	25C34	67CE1	4.011	25C34	670H	4.411
25C34	67CZ	4.629	25C34	160C	4.805	25C34	67CD1	4.850
25C34	160CB	4.898	25C35	67CE1	3.694	25C35	670H	4.238
25C35	209CD2	4.447	25C35	67CZ	4.449	25C35	67CD1	4.551
25C35	1600	4.567	25C35	160CB	4.790	25C35	209CD1	4.791
25N36	1600	4.470	25N36	160CB	4.515	25N36		4.774
25N36	160CG	4.825	25N36	67CE1	4.846	25N36	670H	4.964
25C37	1600	3.691	25C37	160CB	4.458	25C37		4.495
25C37	160C	4.634	25C37	160CA	4.822	25C37	160CG	4.973
25C37	1580	4.994	25N38	1600	3.211	25N38	160C	4.305
25N38	160CB	4.681	25N38	160N	4.817	25N38	160CA	4.898

TABLE XXIX

Active site amino acid residues for Cathepsin K

ASN	18	GLN	19	GLY	20	GLN	21
CYS	22	GLY	23	SER	24	CYS	25
TRP	26	ALA	27	PHE	28	SER	29
GLU	59	ASN	60	ASP	61	GLY	64
GLY	65	GLY	66	TYR	67	MET	68
ASN	70	ALA	134	ALA	137	SER	138
GLN	143	ASP	158	ASN	159	LEU	160
ASN	161	HIS	162	ALA	163	SER	183
TRP	184	TRP	188	LEU	209		

#### WHAT IS CLAIMED IS:

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1. A method of inhibiting cathepsin K which comprises administering to a mammal in need thereof a compound that fits spatially into the active site of cathepsin K, said compound comprising any two of the following:

- (i) an electrophilic carbon atom that binds to the side chain sulfur atom of cysteine 25 wherein said electrophilic carbon atom is 1.7-4.0Å from said sulfur atom:
- (ii) a hydrophobic group that interacts with tryptophan 184 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tryptophan 184 is 4.10-7.10Å;
- (iii) a hydrophobic group that interacts with tyrosine 67, methionine 68, alanine 134, leucine 160, and leucine 209, creating a hydrophobic pocket, and has distance ranges between the centroid of said hydrophobic group and the centroids of the side chain atoms of the amino acid residues of said hydrophobic pocket which are tyrosine 67: 4.91-5.91Å, methionine 68: 5.74-6.74Å, alanine 134: 4.15-5.15Å, leucine 160: 6.18-7.18Å, and leucine 209: 5.71-6.71Å;
- (iv) a hydrophobic group that interacts with tyrosine 67 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tyrosine 67 is 4.10-7.10Å;
- (v) an amino group with a pKa of less than 7 or an oxygen atom, each of which interacts with a hydrogen atom donated by the amide nitrogen of glycine 66 wherein the distance between these two atoms is 2.7-3.5Å;
- (vi) a hydrophobic group that interacts with the main chain atoms of giutamine 21, cysteine 22 and glycine 23 wherein the distance between the centroid of said hydrophobic group and the centroids of glutamine 21, cysteine 22 and glycine 23 are 3.7-5.4, 4.9-5.7 and 5.4-6.7Å, respectively; or
- (vii) a hydrophobic group that interacts with the side chain atoms of glutamine 143 and asparagine 161 and the main chain of alanine 137 and serine 138 wherein the distance between the centroid of the hydrophobic group and the centroids of glutamine 143, asparagine 161, alanine 137, and serine 138 are 7.9-9.6Å, 4.7-5.4Å, 4.2-5.5Å, and 4.6-6.4Å, respectively.

2. A method of inhibiting cathepsin K which comprises administering to a mammal in need thereof a compound that fits spatially into the active site of cathepsin K, said compound comprising any three or more of the following:

(i) an electrophilic carbon atom that binds to the side chain sulfur atom of cysteine 25 wherein said electrophilic carbon atom is 1.7-4.0Å from said sulfur atom:

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- (ii) a hydrophobic group that interacts with tryptophan 184 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tryptophan 184 is 4.10-7.10Å;
- (iii) a hydrophobic group that interacts with tyrosine 67, methionine 68, alanine 134, leucine 160, and leucine 209, creating a hydrophobic pocket, and has distance ranges between the centroid of said hydrophobic group and the centroids of the side chain atoms of the amino acid residues of said hydrophobic pocket which are tyrosine 67: 4.91-5.91Å, methionine 68: 5.74-6.74Å, alanine 134: 4.15-5.15Å, leucine 160: 6.18-7.18Å, and leucine 209: 5.71-6.71Å;
- (iv) a hydrophobic group that interacts with tyrosine 67 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tyrosine 67 is 4.10-7.10Å;
- (v) an amino group with a pKa of less than 7 or an oxygen atom, each of which interacts with a hydrogen atom donated by the amide nitrogen of glycine 66 wherein the distance between these two atoms is 2.7-3.5Å;
- (vi) a hydrophobic group that interacts with the main chain atoms of glutamine 21, cysteine 22 and glycine 23 wherein the distance between the centroid of said hydrophobic group and the centroids of glutamine 21, cysteine 22 and glycine 23 are 3.7-5.4, 4.9-5.7 and 5.4-6.7Å, respectively; or
- (vii) a hydrophobic group that interacts with the side chain atoms of glutamine 143 and asparagine 161 and the main chain of alanine 137 and serine 138 wherein the distance between the centroid of the hydrophobic group and the centroids of glutamine 143, asparagine 161, alanine 137, and serine 138 are 7.9-9.6Å, 4.7-5.4Å, 4.2-5.5Å, and 4.6-6.4Å, respectively.
- 3. A method of inhibiting cathepsin K which comprises administering to a marnmal in need thereof a compound that fits spatially into the active site of cathepsin K, said compound comprising:

(i) an electrophilic carbon atom that binds to the side chain sulfur atom of cysteine 25 wherein said electrophilic carbon atom is 1.7-4.0Å from said sulfur atom; and

- (ii) a hydrophobic group that interacts with tryptophan 184 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tryptophan 184 is 4.10-7.10Å.
  - 4. The method of claim 3 wherein said hydrophobic group that interacts with tryptophan 184 is an aromatic group.

5. The method of claim 4 wherein the centroid of said aromatic group that interacts with tryptophan 184 is 9.24-11.24Å from the centroid of said electrophilic carbon that binds to the side chain sulfur atom of cysteine 25.

- 15 6. The method of claim 3 wherein said electrophilic carbon that binds to the side chain sulfur atom of cysteine 25 is a carbonyl carbon.
  - 7. The method of claim 3 wherein the compound further comprises a hydrophobic group that:

has a centroid which is 5.44-6.94Å from said electrophilic carbon; interacts with tyrosine 67, methionine 68, alanine 134, leucine 160, and leucine 209, creating a hydrophobic pocket; and

has distance ranges between the centroid of said hydrophobic group and the centroids of the side chain atoms of the amino acid residues of said hydrophobic pocket which are tyrosine 67: 4.91-5.91Å, methionine 68: 5.74-6.74Å, alanine 134: 4.15-5.15Å, leucine 160: 6.18-7.18Å, and leucine 209: 5.71-6.71Å.

- 8. The method of claim 7 wherein said hydrophobic group that interacts with said hydrophobic pocket is an isobutyl group.
- 9. The method of claim 3 wherein the compound further comprises a hydrophobic group that interacts with tyrosine 67 wherein the distance between the centroid of said hydrophobic group and the centroid of the side chain atoms of tyrosine 67 is 4.10-7.10Å.

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10. The method of claim 9 wherein said hydrophobic group that interacts with tyrosine 67 is an aromatic group.

- 11. The method of claim 3 wherein the compound further comprises an amino group with a pKa of less than 7 or an oxygen atom, each of which interacts with a hydrogen atom donated by the amide nitrogen of glycine 66 wherein the distance between these two atoms is 2.7-3.5Å.
- 12. The method of claim 3 wherein the compound further comprises a hydrophobic group that interacts with the main chain atoms of glutamine 21, cysteine 22 and glycine 23 wherein the distance between the centroid of said hydrophobic group and the centroids of glutamine 21, cysteine 22 and glycine 23 are 3.7-5.4, 4.9-5.7 and 5.4-6.7Å, respectively.
  - 13. The method of claim 12 wherein said hydrophobic group that interacts with glutamine 21, cysteine 22 and glycine 23 is an isobutyl group.

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- 14. The method of claim 3 wherein the compound further comprises a hydrophobic group that interacts with the side chain atoms of glutamine 143 and asparagine 161 and the main chain of alanine 137 and serine 138 wherein the distance between the centroid of the hydrophobic group and the centroids of glutamine 143, asparagine 161, alanine 137, and serine 138 are 7.9-9.6Å, 4.7-5.4Å, 4.2-5.5Å, and 4.6-6.4Å, respectively.
- 15. The method of claim 1 wherein the compound is:

  3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone;

4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyπolidinone;

4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-N-[N-(methyl)-L-leucyl)]-3-pyrrolidinone;

4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone; bis-(Cbz-leucinyl)-1,3-diamino-propan-2-one;

2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-leucinyl)]carbohydrazide;

- (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide;
- 1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one; or
- 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide; or a pharmaceutically acceptable salt thereof.
- 16. A composition comprising cathepsin K in crystalline form.
- 17. The composition according to claim 16 wherein cathepsin K has an active site cavity formed by the amino acids in Table XXIX.
- 15 18. The composition of claim 17 wherein said active site is characterized by the coordinates selected from the group consisting of the coordinates of Tables I-X.
  - 19. A cathepsin K crystal.

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20. An isolated, properly folded cathepsin K molecule or fragment thereof having a conformation comprising a catalytically active site formed by the residues listed in Table XXIX, said active site defined by the protein coordinates of Table I.

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- 21. A peptide, peptidomimetic or synthetic molecule which binds with the active site cavity of cathepsin K according to claim 17.
- 22. A method of identifying an inhibitor compound capable of binding to,
  and inhibiting the proteolytic activity of, cathepsin K, said method comprising:
  introducing into a suitable computer program information defining an
  active site conformation of a cathepsin K molecule comprising a catalytically active
  site formed by the residues listed in Table XXIX, said active site defined by the
  protein coordinates of Table I, wherein said program displays the three-dimensional
  structure thereof:

creating a three dimensional representation of the active site cavity in said computer program;

displaying and superimposing the model of said test compound on the model of said active site;

assessing whether said test compound model fits spatially into the active site;

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preparing said test compound that fits spatially into the active site; using said test compound in a biological assay for a protease characterized by said active site; and

determining whether said test compound inhibits cathepsin K activity in said assay.

- 23. A peptide, peptidomimetic or synthetic molecule identified by the method of Claim 22.
- 24. A method of drug design comprising using the structural coordinates of a cathepsin K crystal to computationally evaluate a chemical entity for associating with the active site of cathepsin K.
- 25. The method according to claim 24, wherein said entity is a competitive or non-competitive inhibitor of cathepsin K.
- 26. A method for identifying inhibitors which competitively bind to the active site of a cathepsin K molecule or fragment thereof characterized by a catalytically active site formed by the residues listed in Table XXIX, said method comprising the steps of:

providing the coordinates of said active site of the protease to a computerized modeling system;

identifying compounds which will bind to the structure; and screening the compounds identified for protease inhibitory bioactivity.

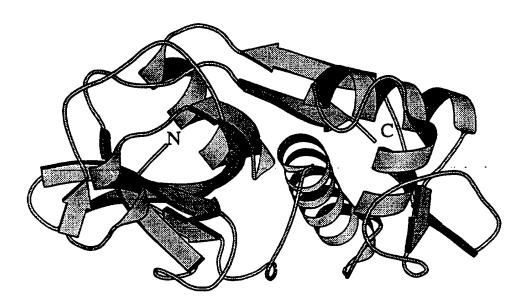
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# FIGURE 1 Sequence Comparison Between Cathepsin K and the Papain Superfamily of Cysteine Proteases

	1				50				
CALE		. MWGLKVLLL	PVVSFA	LYPEEILDT	HWELWICKTHR	KQ.YNNKVDE	ISRRLIWENN	LKYISIHNLE	ASLGVHTYEL
Cats		MCRLVCVLLV	CSSAVA	QLHKDPTLDH	HWHLMKKTYG	KQ.YKEKWEE	AVRRLIWEEN	LKPVMLHNLE	HSMGNHSYDL
Catl		MNPTLIL	AAPCLGIASA	TLTFDHSLEA	OMLKMKYRODI	RL.Y.GMMEE	GWRRAVWERN	MONI ELHNOE	YREGICHSFTM
Papain	• • • • • • • • • • • • • • • • • • • •	•••••		• • • • • • • • • • • • • • • • • • • •	• • • • • • • • • • • • • • • • • • • •	• • • • • • • • • • • • • • • • • • • •	•••••	• • • • • • • • • • • • • • • • • • • •	
lotinidin	MGLPKSFVSH	SLLPPSTLLI	LSLAFNAKNL	TORTNDEVKA	Myeswlikyg	ks . Ynslgew	ERRFEIFKET	LRFIDENNAD	TNRSYKV
Cata	MA	TLPLLCAGAW	LLGVPVCGAA	elsvnslekf	HPKSWMSKHR	KT.Y.STEEY	HHRLQTPASN	WRKINAHN	NGNHTFKM
Cata		• • • • • • • • • • • • • • • • • • • •		• • • • • • • • • • • • • • • • • • • •	MWQLWASLCC	LLVLANARSR	PSPHPVSDEL	VNYVNKRNTT	wqaghnfynv
	100					150			
Cati	AMPHICOMTS	EEVVQXXXTGL	KVPLSHSRSN	DTLYIPEWEG	RAPDSVDYRK	KG.YVTPVKN	OGOCGSCHAF	SSVGALEGQL	KKKTGKLLN.
Cats	CHANHLEDMYS	EEVMSLTSSL	RVP.SQWQRN	IT.YKSNPNR	ILPDSVDWRE	KG. CVTEVKY	QGSCGACHAP	SAVGALEAQL	KLKTGKLVT.
Catl	amnafgements	eefrqvioigf	QNRKPRK	GKVFQEPLTY	Eaprsvowre	KG.YVTPVKN	QGQCGSCWAP	SATGALEGOM	PRETGRLIS.
Papain					. IPEYVDWRQ	KG. AVTPVKN	QGSCGSCWAP	SAVVTIEGII	Kirtgning.
ctinidin	GLNQFADLTD	EEFRSTYLGF	. TSGSNKTKV	SNRYEPRFGQ	VLPSYVDWRS	AG.AVVDIKS	QGBCGGCNAP	SALATVEGIN	KIVIGVLIS.
Cati	Alnopsimsp	ABIKHKY	LWSEPQNCSA	TKSNYLRGTG	PYPPSVDWRK	KGNFVSPVKN	QGACGSCHTP	STTGALESAI	AIATGROULS.
Cata	DMSYLKRLCG	TFLGGPKPPQ	RVMFTEDLKL	PASPDAR	EQWP	OCPTIKEIRD	QGSCGSCHAF	GAVEAISDRI	CIHTNAHVSV
		200					250		
Catx	.LSPQNLVDC	VSENDGC	GGGYMTNAFQ	YVQXXXGIDS	EDAY		PYV	GQEESCH	YNPTG
Cats	.LSAQNLVDC	STERYGNIKGC	NGGFMTTAFQ	YIIDNKGIDS	DASY			AMDQKCQ	YDSKY
Catl	. LSEQNLVDC	SGPQ.GNEGC	NGGLMDYAFQ	YVQDNGGLDS	eesy		PYE	ATERSCK	YNPKY
Papain	.YSEQELLDC	DRRSY.GC	NGGYPWSALQ	LVAQY.GIHY	RNTY			GVQRYCR	SREKG
ctinidia	.LSEQELIDC	GRTQNTR.GC	NGGYITDGFQ	FIINNGGINT	<b>EENY</b>		PYT	AQDGECN	LDLQN
Cate	.LAEQQLVDC	A. QDFRRYGC	QGGLPSQAFE	YILYNKGIMG	EDTY			GKDGYCK	FQP.G
Cath	EVSAEDLLTC	CGSMCG.DGC	NGGYPAZAWN	P.WTRKGLVS	GGLYESHVGC	RPYSIPPCEH	HVNGSRPPCT	GEGDTPKCSK	ICEPGYSPTY
			300					350	
Cati	K.AAKCRGYR	EIPEGNEKAL	KRAVAFVGPV	SVAIDASLTS	POPVSKG <b>VY</b> Y	DESC. NEDN	LNHAVLAVGY	GIQKGN	XIMIIKISWG
Cats	R.AATCSKYT	ELPYGREDVL	KZKVANKGPV	SVGVDARHPS	FFLYRSGVYY	epscTon	VNHGVLVVGY	GDLNGK	EYWLVKNSWG
CatL	S.VANDTGPV	DIP.RQEKAL	HIKAVATVGPI	SVAIDAGHES	FLEYNDRIYE	EPICSSED	HOHGVLVVGY	GPESTESCHN	KYWLVKNSWG
Papain	PYAAKTDGVR	QVQPYNQGAL	LYSIAN.QPV	SVVLQAAGND	POLYRGGIFV	GPCGNK	VDHAVAAVGY	GP	nyiliknswg
ctinidin	EXYVTIDIYE	NVPYNNEWAL	QTAVTY.QPV	SVALDAAGDA	FXHYSSGIFT	GPCGTA	IDHAVTIVGY	GTEGGI	DYWIVKNSWD
Cath	KAIGFVKDVA	NITIYDEEAM	VEXVALYMPV	SPAPEVTCO.	FMM:RTGIYS	STSCHKTPDK	VNHAVLAVCY	GEXXIGI	PYWIVIONSWG
CatB	KODIOHYGYNS	YSVSNSEKDI	MAEIYKNGPV	EGAFSV.YSD	PLLYKSGVYQ	HVTGERRIG	.GHAIRILGW	GVENGT	PYWLVANSWN
				400			430		
Cati	ENWGNKGYIL	MARNIKONA	. CGIANLASP	PXM					
Cats	HNFGEEGYIR	MARNIKGNH	.CGIASPPSY	PE1					
Catl	EEWGMGGYVK	MAKDRRNH	. CGIASAASY	PTV					
Papain	TCWGENGYIR	IKRGTONSYG	VCGLYTSSFY	PV:0N					
ctinidin	TTWGEEGYMR	ILRNVGGA.G	TCGIATMPSY	PVKYNNONAP	KPYSSLINPP	apsmskogpv	GVDDGQRYSA		
		TERCEN							

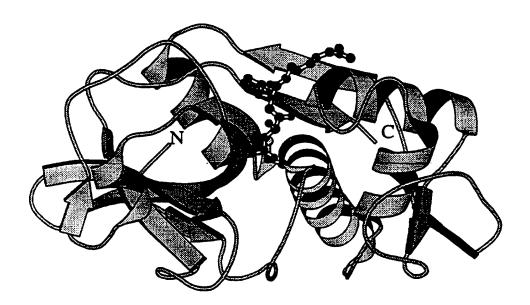
Cats TDWGDNGFFK ILRGQDHCGI ESEVVAGIPR TDQYWEKI......

# FIGURE 2



Human Cathepsin K

## FIGURE 3

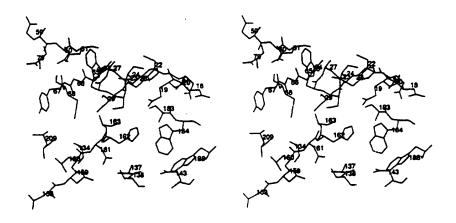


Human Cathepsin K E-64

Figure 4a

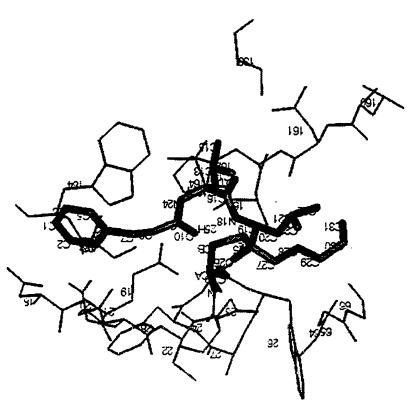
Cathepsin K Active Site

Figure 4b



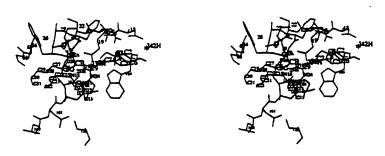
Stereo View Cathepsin K Active Site

## FIGURE 5a



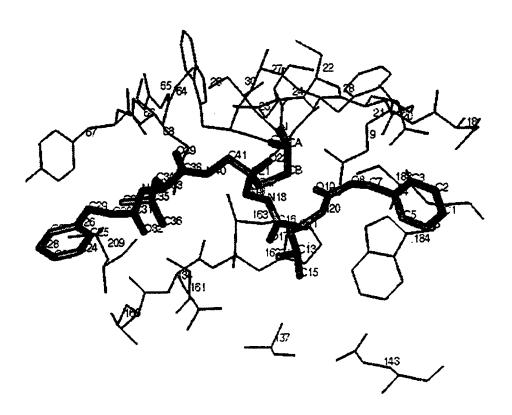
Inhibitor = 3(3)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone

## FIGURE 5b



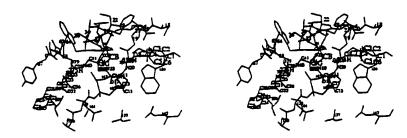
Inhibitor = 3(S)-3-[(N-benzyloxycarbonyl)-L-leucinyl]amino-5-methyl-1-(1-propoxy)-2-hexanone

## FIGURE 6a



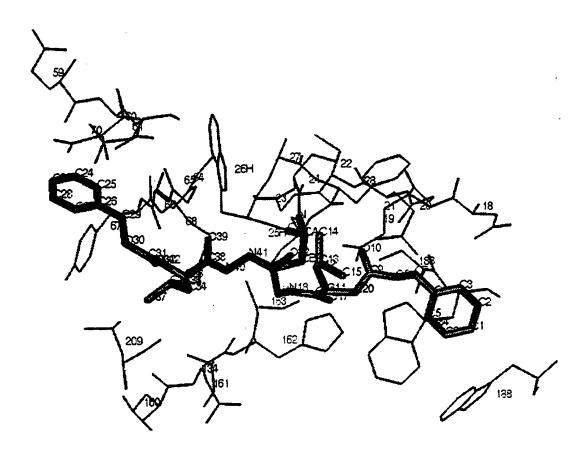
Inhibitor = bis-(cbz-leucinyl)-1,3-diamino-propan-2-one

## FIGURE 6b



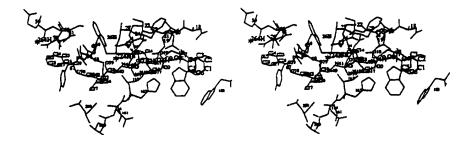
Inhibitor = bis-(cbz-leucinyl)-1,3-diamino-propan-2-one

### FIGURE 7a



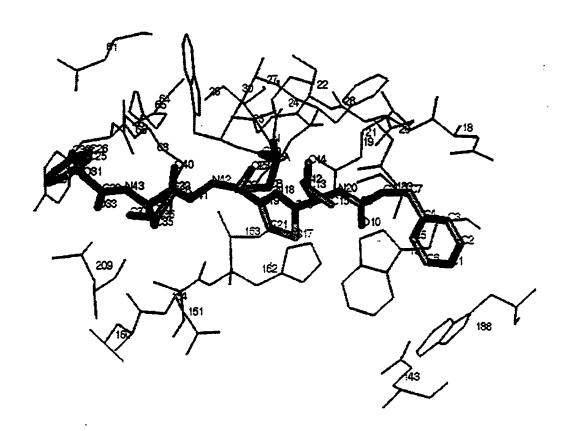
 $Inhibitor = 2.2 \text{--} N, N'\text{--}bis\text{--}benzyloxycarbonyl-L--leucinylcarbohydrazide}$ 

## FIGURE 7b



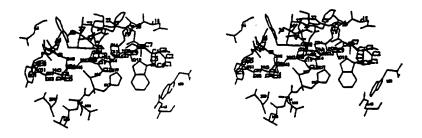
Inhibitor = 2,2'-N,N'-bis-benzyloxycarbonyl-L-leucinylcarbohydrazide

## FIGURE 8a



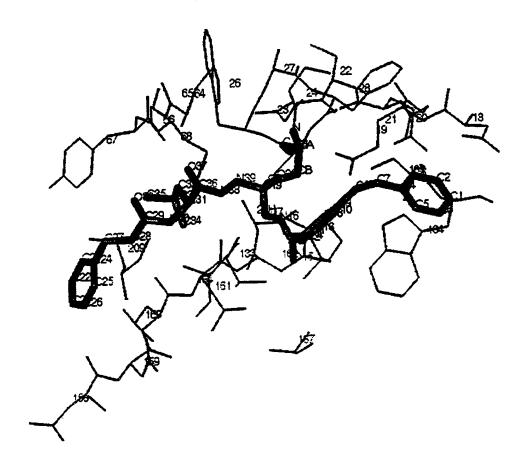
Inhibitor = (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide

### FIGURE 8b



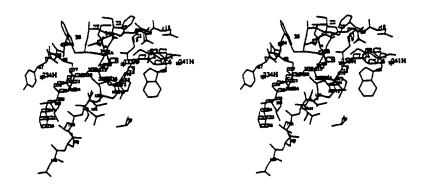
Inhibitor = (1S)-N-[2-[(1-benzyloxycarbonylamino)-3-methylbutyl]thiazol-4-ylcarbonyl]-N'-(N-benzyloxycarbonyl-L-leucinyl)hydrazide

## FIGURE 9a



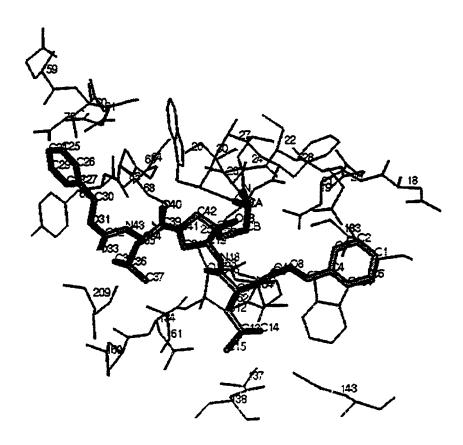
Inhibitor = 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxycarbonyl-L-benzyloxycarbohydrazide)] carbohydrazide

### FIGURE 9b



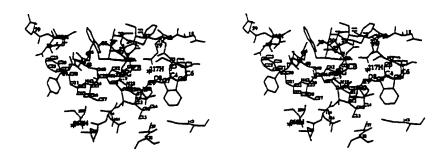
 $\label{eq:loss_entropy} Inhibitor = 2-[N-(3-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxybenzoyl-L-benzyloxybenzoyl-L-benzyloxybenzoyl)]-2'-[N'-(N-benzyloxybenzoyl-L-benzyloxyb$ 

### FIGURE 10a



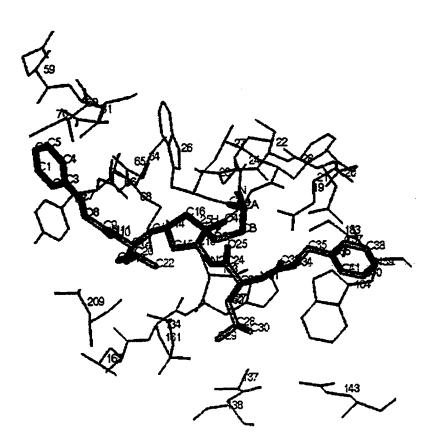
Inhibitor = 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

### FIGURE 10b



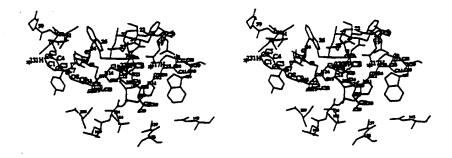
Inhibitor = 4-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

## FIGURE 11a



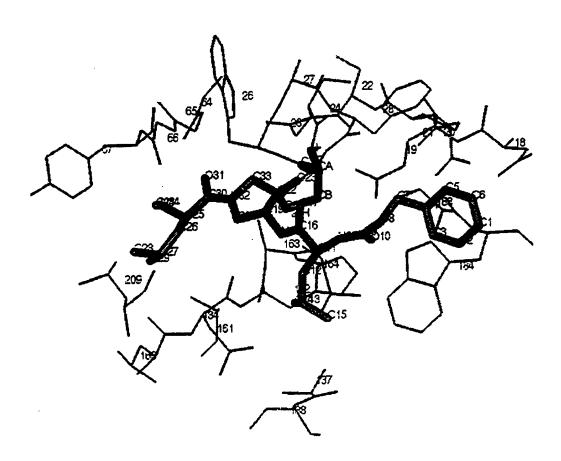
Inhibitor = 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

## FIGURE 11b



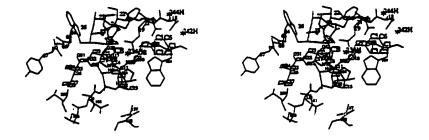
Inhibitor = 4-[N-[(4-pyridylmethoxy)carbonyl]-L-leucyl]-1-[N-[(phenylmethoxy)carbonyl]-L-leucyl]-3-pyrrolidinone

## FIGURE 12a



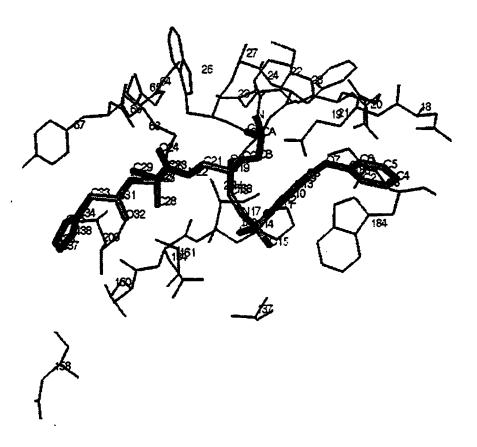
Inhibitor = 4-[N-[(phenylmethoxy)carbony!]-L-leucyl]-1-N[N-(methyl)-L-leucyl]-3-pyrrolidinone

## FIGURE 12b



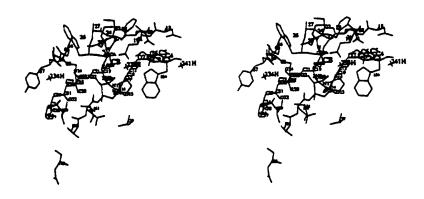
Inhibitor = 4-[N-[(phenylmethoxy)carbonyi]-L-leucyl]-1-N[N-(methyl)-L-leucyl]-3-pyrrolidinone

## FIGURE 13a



Inhibitor =1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one

### FIGURE 13b



Inhibitor = 1-N-(N-imidazole acetyl-leucinyl)-amino-3-N-(4-phenoxy-phenyl-sulfonyl)-amino-propan-2-one

#### INTERNATIONAL SEARCH REPORT

International application No. PCT/US96/17512

A. CLA	SSIFICATION OF SUBJECT MATTER			
	Please See Extra Sheet.	617		
According to	435/23, 24, 212, 226; 514/19, 365, 370, 400, 615, o International Patent Classification (IPC) or to both	national classification and IPC		
B. FIEL	DS SEARCHED			
Minimum d	ocumentation searched (classification system followed	d by classification symbols)		
	435/23, 24, 212, 226; 514/19, 365, 370, 400, 615, 6			
Documentat	ion searched other than minimum documentation to the	e extent that such documents are included	in the fields searched	
Electronic d	ata base consulted during the international search (na	ime of data base and, where practicable,	, search terms used)	
APS, DIA search te	NLOG erms: cathepsin, osteoclast, inhibit, crystal, leu	cine, thiazol		
C. DOC	UMENTS CONSIDERED TO BE RELEVANT		<b>:</b>	
Category*	Citation of document, with indication, where ap	propriate, of the relevant passages	Relevant to claim No.	
Y			16-20	
Y, P		22-26 20,807 A (LAVIN ET AL) 19 March 1996 (19/03/96), 7, lines 12-36, column 9, lines 1-56. 31,573 A (BALAJI ET AL) 19 July 1994 (19/07/94), 8, line 1 - column 9, line 63. 301,969 A (HASTINGS ET AL) 26 March 1996 1-14, 16-26		
Y	US 5,331,573 A (BALAJI ET AL) column 8, line 1 - column 9, line 6		22-26	
Y, P				
X Furth	er decuments are listed in the continuation of Box C		· · · · · · · · · · · · · · · · · · ·	
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the	priority data claimed actual completion of the international search	Date of mailing of the international sec	irch report	
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International application No.
PCT/US96/17512

Category*	Citation of document, with indication, where appropriate, of the relevant passages	passages Relevant to claim No	
X  Y	US 5,424,325 A (ANDO ET AL) 13 June 1995 (13/06/95), column 1, lines 5-9, column 2, line 62 - column 3, line 5, column 4, lines 6-53.	21,23  1-14	
X  Y	US 5,422,359 A (ANDO ET AL) 06 June 1995 (06/06/95), column 1, lines 5-9, column 2, line 62 - column 3, line 5, column 4, lines 7-52.	21,23  1-14	
х — Y	US 5,223,486 A (GORDON ET AL) 29 June 1993 (29/06/93), column 3, lines 26-44, column 4, lines 36-42.	21,23	
х — Y	US 5,395,824 A (HIGUCHI ET AL) 07 March 1995 (07/03/95), column 2, line 1 - column 3, line 14.	21,23	
A, P	BOSSARD et al. Proteolytic Activity of Human Osteoclast Cathepsin K. The Journal Of Biological Chemistry. 24 May 1996, Volume 271, Number 21, pages 12517-12524.	1-26	
Y	DESJARLAIS et al. Using Shape Complementarity as an Initial Screen in Designing Ligands for a Receptor Binding Site of Known Three-Dimensional Structure. Journal of Medicinal Chemistry. 1988, Volume 31, Number 4, pages 722-729, especially the abstract.	22-26	
X, P  Y, P	BROMME et al. Peptidyl vinyl sulphones: a new class of potent and selective cysteine protease inhibitors. Biochemical Journal. 1996, Volume 315, pages 85-89, especially the abstract, Figure 1.	21,23	
<u>х</u> <u>т</u>	VELASCO et al. Human Cathepsin O. Molecular Cloning From a Breast Carcinoma, Production Of the Active Enzyme In Escherichia Coli, And Expression Analysis In Human Tissues. The Journal Of Biological Chemistry. 28 October 1994, Volume 269, Number 43, pages 27136-27142, especially the abstract.	21,23 16-20	
<u>х</u> <u>-</u> Y	MAGRATH et al. Cysteine Protease Inhibition by Azapeptide Esters. Journal Of Medicinal Chemistry. 1992, Volume 35, Number 23, pages 4279-4283, especially page 4281, column 1, structures 1-4 and 7.	21, 23	
X Y	GRAYBILL et al. Synthesis And Evaluation Of Azapeptide- Derived Inhibitors Of Serine And Cysteine Proteases. Bioorganic & Medicinal Chemistry Letters. 1992, Volume 2, Number 11, pages 1375-1380, especially page 1377, Scheme I.	21, 23	

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International application No. PCT/US96/17512

A61K 31/16, 31/165,	31/415, 31/425, 38/05; C12	2N 9/48, 9/64; C12Q	1/37	
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